



## SEQUENCE LISTING

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<120> Proteins

<130> PWC/P21089wo

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<151> 1999-03-19

<160> 212

<170> PatentIn Ver. 2.1

<210> 1

<211> 1248

<212> DNA

<213> Streptococcus agalactiae

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<210> 2

<211> 415

<212> PRT

<213> Streptococcus agalactiae

<400> 2

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1 5 10 15

Ser Val Val Ala Gly Gly Ala Ile Ala Ala Thr His Ser Asn Ser Val 20 25 30

Asp Ala Ala Ser Lys Lys Thr Ile Lys Leu Trp Val Pro Thr Asp Ser 35 40 45

Lys Ala Ser Tyr Lys Ala Ile Val Lys Lys Phe Glu Lys Glu Asn Lys
50 55 60

Gly Val Thr Val Lys Met Ile Glu Ser Asn Asp Ser Lys Ala Gln Glu
65 70 75 80



Asn Val Lys Lys Asp Pro Ser Lys Ala Ala Asp Val Phe Ser Leu Pro 85 90 95

His Asp Gln Leu Gly Gln Leu Val Glu Ser Gly Val Ile Gln Glu Ile
100 105 110

Pro Glu Gln Tyr Ser Lys Glu Ile Ala Lys Asn Asp Thr Lys Gln Ser 115 120 125

Leu Thr Gly Ala Gln Tyr Lys Gly Lys Thr Tyr Ala Phe Pro Phe Gly
130 135 140

Asp Val Lys Ser Tyr Glu Thr Ile Thr Ser Lys Gly Lys Phe Gly Gln
165 170 175

Gln Leu Lys Ala Ala Asn Ser Tyr Val Thr Gly Pro Leu Phe Leu Ser 180 185 190

Val Gly Asp Thr Leu Phe Gly Lys Ser Gly Glu Asp Ala Lys Gly Thr
195 200 205

Asn Trp Gly Asn Glu Ala Gly Val Ser Val Leu Lys Trp Ile Ala Asp 210 215 220

Gln Lys Lys Asn Asp Gly Phe Val Asn Leu Thr Ala Glu Asn Thr Met 225 230 235 240

Ser Lys Phe Gly Asp Gly Ser Val His Ala Phe Glu Ser Gly Pro Trp
245 250 255

Asp Tyr Asp Ala Ala Lys Lys Ala Val Gly Glu Asp Lys Ile Gly Val
260 265 270

Ala Val Tyr Pro Thr Met Lys Ile Gly Asp Lys Glu Val Gln Gln Lys 275 280 285



Ala Phe Leu Gly Val Lys Leu Tyr Ala Val Asn Gln Ala Pro Ala Gly 290 295 300

Ser Asn Thr Lys Arg Ile Ser Ala Ser Tyr Lys Leu Ala Ala Tyr Leu 305 310 315 320

Thr Asn Ala Glu Ser Gln Lys Ile Gln Phe Glu Lys Arg His Ile Val

Pro Ala Asn Ser Ser Ile Gln Ser Ser Asp Ser Val Gln Lys Asp Glu 340 345 350

Leu Ala Lys Ala Val Ile Glu Met Gly Ser Ser Asp Lys Tyr Thr Thr 355 360 365

Val Met Pro Lys Leu Ser Gln Met Ser Thr Phe Trp Thr Glu Ser Ala 370 375 380

Ala Ile Leu Ser Asp Thr Tyr Ser Gly Lys Ile Lys Ser Ser Asp Tyr 385 390 395 400

Leu Lys Arg Leu Lys Gln Phe Asp Lys Asp Ile Ala Lys Thr Lys
405 410 415

<210> 3

<211> 1539

<212> DNA

<213> Streptococcus agalactiae

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aacagtagtc aagcagacag taagccaggt caatcaacaa agactgaatt aaaacctgag 420 cctaccttac cattagtaga gcctaaaata actcccgctc cgtctcagat agaaagtgtt 480 cagacaaatc agaatgcttc tgttcctgct ttatcctttg atgataactt attatcaaca 540 ccqatttcac cagtgacagc aacgccattc tacgtagaac actggtctgg tcaggatgcc 600 tactctcact atttattgtc acatcgttac ggtatcaaag ctgaacaatt agatgggtac 660 ttaaaatctt tagggattca atatgattct aatcgtatca atggtgctaa gttattacaa 720 tgggaaaaag atagtggttt agatgtccgt gctattgtag ctattgctgt ccttgaaagt 780 tcattgggaa ctcaaggagt ggctaaaatg ccaggtgcta atatgtttgg ttatggtgcc 840 tttgatcatg actctagcca tgctagtgct tataatgatg aagaagcaat tatgttgttg 900 acaaaaaata caattattaa aaacaacaac tctagctttg aaatccaaga tttgaaagca 960 cagaaattat cttctggaca acttaataca gttactgagg gtggtgttta ttatacagat 1020 aactctggaa ctggtaaacg tcgtgcccag attatggaag atttagaccg ctggattgat 1080 caacatggag ggacaccaga aattcctgct gccttgaaag ctttatcgac agcaagttta 1140 gcagatttac caagtggttt tagcttatca acagcggtta acacagctag ctatattgca 1200 tcaacttatc catggggtga atgtacatgg tatgtcttta accgcgctaa agagttaggt 1260 tatacatttg atccatttat gggtaatggt ggagattggc aacataaggc tggctttgaa 1320 acaacacatt caccaaaagt aggctatgct gtatcatttt caccaggaca agctggtgct 1380 gatggcactt acggtcacgt agctattgtt gaagaagtta aaaaagatgg ttcagttctc 1440 atttcagaat ctaatgcaat gggacgtggt attgtctctt accgtacttt tagttcagca 1500

1539

<210> 4

<211> 512

<212> PRT

<213> Streptococcus agalactiae

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<400> 4

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Leu Ser Leu Ser Ser Pro Leu Val Thr Leu Ala Glu Thr Ile Asn Pro
20 25 30

Glu Thr Ser Leu Thr Met Ala Thr Ala Ser Thr Glu Ser Ser Glu
35 40 45



Ala Glu Lys Gln Glu Lys Thr Gln Pro Thr Asp Ser Glu Thr Ala Ser 

Pro Ser Ala Glu Gly Ser Ile Ser Thr Glu Lys Thr Glu Ile Gly Thr 

Thr Glu Thr Ser Ser Ser Asn Glu Ser Ser Ser Ser Ser His Gln 

Ser Ser Ser Asn Glu Asp Ala Lys Thr Ser Asp Ser Ala Ser Thr Ala 

Ser Thr Pro Ser Thr Asn Thr Thr Asn Ser Ser Gln Ala Asp Ser Lys 

Pro Gly Gln Ser Thr Lys Thr Glu Leu Lys Pro Glu Pro Thr Leu Pro 

Leu Val Glu Pro Lys Ile Thr Pro Ala Pro Ser Gln Ile Glu Ser Val 

Gln Thr Asn Gln Asn Ala Ser Val Pro Ala Leu Ser Phe Asp Asn 

Leu Leu Ser Thr Pro Ile Ser Pro Val Thr Ala Thr Pro Phe Tyr Val 

Glu His Trp Ser Gly Gln Asp Ala Tyr Ser His Tyr Leu Leu Ser His 

Arg Tyr Gly Ile Lys Ala Glu Gln Leu Asp Gly Tyr Leu Lys Ser Leu 

Gly Ile Gln Tyr Asp Ser Asn Arg Ile Asn Gly Ala Lys Leu Leu Gln 

Trp Glu Lys Asp Ser Gly Leu Asp Val Arg Ala Ile Val Ala Ile Ala 



Val Leu Glu Ser Ser Leu Gly Thr Gln Gly Val Ala Lys Met Pro Gly
260 265 270

Ala Asn Met Phe Gly Tyr Gly Ala Phe Asp His Asp Ser Ser His Ala 275 280 285

Ser Ala Tyr Asn Asp Glu Glu Ala Ile Met Leu Leu Thr Lys Asn Thr 290 295 300

Ile Ile Lys Asn Asn Asn Ser Ser Phe Glu Ile Gln Asp Leu Lys Ala 305 310 315 320

Gln Lys Leu Ser Ser Gly Gln Leu Asn Thr Val Thr Glu Gly Gly Val
325 330 335

Tyr Tyr Thr Asp Asn Ser Gly Thr Gly Lys Arg Arg Ala Gln Ile Met 340 345 350

Glu Asp Leu Asp Arg Trp Ile Asp Gln His Gly Gly Thr Pro Glu Ile 355 360 365

Pro Ala Ala Leu Lys Ala Leu Ser Thr Ala Ser Leu Ala Asp Leu Pro 370 375 380

Ser Gly Phe Ser Leu Ser Thr Ala Val Asn Thr Ala Ser Tyr Ile Ala 385 390 395 400

Ser Thr Tyr Pro Trp Gly Glu Cys Thr Trp Tyr Val Phe Asn Arg Ala 405 410 415

Lys Glu Leu Gly Tyr Thr Phe Asp Pro Phe Met Gly Asn Gly Gly Asp 420 425 430

Trp Gln His Lys Ala Gly Phe Glu Thr Thr His Ser Pro Lys Val Gly
435 440 445

Tyr Ala Val Ser Phe Ser Pro Gly Gln Ala Gly Ala Asp Gly Thr Tyr 450 455 460



470

Gly His Val Ala Ile Val Glu Glu Val Lys Lys Asp Gly Ser Val Leu

480

Ile Ser Glu Ser Asn Ala Met Gly Arg Gly Ile Val Ser Tyr Arg Thr
485 490 495

Phe Ser Ser Ala Gln Ala Ala Gln Leu Thr Tyr Gly Ile Gly His Lys
500 505 510

<210> 5

465

<211> 1293

<212> DNA

<213> Streptococcus agalactiae

<400> 5

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gagaatctta aaacgaatcg tcaagggaaa tcgactatta tttcagcaca tcgtttatca 1140 gctgttgtgc acgcagacct tatcttagtt atgcgagacg gcagagtcat tgagcgaggt 1200 caacatcaag agttgctaaa taaaggtggt tggtatgctg aaacgtatgc ctcacagcaa 1260 ttagaaatgg aggaagcatt tgatgaagtc taa 1293

<210> 6

<211> 430

<212> PRT

<213> Streptococcus agalactiae

<400> 6

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1 5 10 15

Ile Ala Leu Gly Cys Met Arg Met Ser Asp Leu Lys Gly Lys Gln Ala
20 25 30

Glu Glu Val Val Gly Thr Ala Leu Asp Leu Gly Ile Ile Asn Asn Lys
35 40 45

Val Gln Glu Ser Val Ser Gly Val Lys Val Thr Lys Ser Leu Cys Tyr 50 55 60

Gln Glu Gln Glu Ile Ala Ser Phe Gln Glu Ile Asn Gln Met Thr Phe
65 70 75 80

Val Lys Asn Met Arg Thr Met Thr Tyr Asp Val Met Phe Asp Pro Leu 85 90 95

Val Leu Leu Phe Ile Gly Ala Ser Tyr Val Leu Thr Leu Ala Met Gly
100 105 110

Ala Phe Met Ile Ser Lys Gly Gln Val Thr Val Gly Asp Leu Val Thr
115 120 125

Phe Val Thr Tyr Leu Asp Met Leu Val Trp Pro Leu Met Ala Ile Gly
130 135 140



Phe Leu Phe Asn Met Val Gln Arg Gly Ser Val Ser Tyr Asn Arg Ile
145 150 155 160

Asn Ser Leu Leu Glu Gln Glu Ser Asp Ile Thr Asp Pro Leu Asn Pro 165 170 175

Ile Lys Pro Val Val Asn Gly Thr Leu Arg Tyr Asp Ile Asp Phe Phe 180 185 190

Arg Tyr Asp Asn Glu Glu Thr Leu Ala Asp Ile His Phe Thr Leu Glu 195 200 205

Lys Gly Gln Thr Leu Gly Leu Val Gly Gln Thr Gly Ser Gly Lys Thr 210 215 220

Ser Leu Ile Lys Leu Leu Leu Arg Glu His Asp Val Thr Gln Gly Lys
225 230 235 240

Ile Thr Leu Asn Lys His Asp Ile Arg Asp Tyr Arg Leu Ser Glu Leu 245 250 255

Arg Gln Leu Ile Gly Tyr Val Pro Gln Asp Gln Phe Leu Phe Ala Thr
260 265 270

Ser Ile Leu Glu Asn Val Arg Phe Gly Asn Pro Thr Leu Ser Ile Asn 275 280 285

Ala Val Lys Glu Ala Thr Lys Leu Ala His Val Tyr Asp Asp Ile Glu 290 295 300

Gln Met Pro Ala Gly Phe Glu Thr Leu Ile Gly Glu Lys Gly Val Ser 305 310 315 320

Leu Ser Gly Gly Gln Lys Gln Arg Ile Ala Met Ser Arg Ala Met Ile 325 330 335

Leu Asp Pro Asp Ile Leu Ile Leu Asp Asp Ser Leu Ser Ala Val Asp
340 345 350



Ala Lys Thr Glu His Ala Ile Val Glu Asn Leu Lys Thr Asn Arg Gln 355 360 365

Gly Lys Ser Thr Ile Ile Ser Ala His Arg Leu Ser Ala Val Val His 370 375 380

Ala Asp Leu Ile Leu Val Met Arg Asp Gly Arg Val Ile Glu Arg Gly 385 390 395 400

Gln His Gln Glu Leu Leu Asn Lys Gly Gly Trp Tyr Ala Glu Thr Tyr
405 410 415

Ala Ser Gln Gln Leu Glu Met Glu Glu Ala Phe Asp Glu Val 420 425 430

<210> 7

<211> 999

<212> DNA

<213> Streptococcus agalactiae

#### <400> 7

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aatttttcaa ccttacaaac atcaatggta tcagcagggc gtgtgtttga tctgattgat 960 gaaacaggtt ttgaaccaag ccaaaaaaat acagaagct 999

<210> 8

<211> 332

<212> PRT

<213> Streptococcus agalactiae

<400> 8

Met Lys Ser Asn Gln Trp Gln Val Phe Lys Arg Leu Ile Ser Tyr Leu 1 5 10 15

Arg Pro Tyr Lys Trp Phe Thr Val Leu Ala Leu Ser Leu Leu Leu Leu 20 25 30

Thr Thr Val Val Lys Asn Ile Ile Pro Leu Ile Ala Ser His Phe Ile
35 40 45

Asp His Tyr Leu Thr Asn Val Asn Gln Thr Ala Val Leu Ile Leu Val 50 55 60

Gly Tyr Tyr Ser Met Tyr Val Leu Gln Thr Leu Ile Gln Tyr Phe Gly
65 70 75 80

Asn Leu Phe Phe Ala Arg Val Ser Tyr Ser Ile Val Arg Asp Ile Arg 85 90 95

Arg Asp Ala Phe Ala Asn Met Glu Arg Leu Gly Met Ser Tyr Phe Asp 100 105 110

Arg Thr Pro Ala Gly Ser Ile Val Ser Arg Ile Thr Asn Asp Thr Glu 115 120 125

Ala Ile Ser Asp Met Phe Ser Gly Ile Leu Ser Ser Phe Ile Ser Ala 130 135 140



Ile Phe Ile Phe Thr Val Thr Leu Tyr Thr Met Leu Met Leu Asp Ile Lys Leu Thr Gly Leu Val Ala Leu Leu Leu Pro Val Ile Phe Ile Leu Val Asn Val Tyr Arg Lys Lys Ser Val Thr Val Ile Ala Lys Thr Arg Ser Leu Leu Ser Asp Ile Asn Ser Lys Leu Ser Gly Ser Ile Glu Gly Ile Arg Ile Val Gln Ala Phe Gly Gln Glu Glu Arg Leu Lys Thr Glu Phe Glu Glu Ile Asn Lys Glu His Val Val Tyr Ala Asn Arg Ser Met Ala Leu Asp Ser Leu Phe Leu Arg Pro Ala Met Ser Leu Leu Lys Leu Leu Ala Tyr Ala Val Leu Met Ser Tyr Phe Gly Phe Thr Gly Val Lys Gly Gly Leu Thr Ala Gly Leu Met Tyr Ala Phe Ile Gln Tyr Val Asn Arg Leu Phe Asp Pro Leu Ile Glu Val Thr Gln Asn Phe Ser Thr Leu Gln Thr Ser Met Val Ser Ala Gly Arg Val Phe Asp Leu Ile Asp Glu - 320

Thr Gly Phe Glu Pro Ser Gln Lys Asn Thr Glu Ala 325 330







<211> 3753

<212> DNA

<213> Streptococcus agalactiae

<400> 9

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gcagcctttt cagagaaact agattattta cagaaattag gagttacaca cattcagctt 1980 ttaccggtat tgagttattt ttatgttaat gaaatggata agtcacgctc aacagcttac 2040 acttcctcag acaataatta caattggggc tatgacccac agagctattt tgctctttct 2100 ggaatgtatt cagagaaacc aaaagatcca tcagcacgta tcgccgaatt aaaacaatta 2160 atacatgata ttcataaacg tggcatgggg gttatacttg atgtcgtcta taatcacact 2220 gcaaaaactt atctctttga ggatatagaa cctaattatt atcactttat gaatgaagat 2280 ggttcaccaa gagaaagttt tggaggggga cgtttaggaa ccactcatgc aatgagtcgt 2340 cgtgttttgg ttgattccat taaatatctt acaagtgaat ttaaagttga tggtttccgt 2400 tttgatatga tgggagatca tgatgcggct gcgattgaat tagcttataa agaagctaaa 2460 gctattaatc ctaatatgat tatgattggt gagggctgga gaacattcca aggcgatcaa 2520 ggtaagccgg ttaaaccagc tgaccaagat tggatgaagt caaccgatac agttggcgtc 2580 ttttcagatg atattcgtaa tagcttgaaa tctggttttc caaatgaagg tactccagct 2640 ttcatcacag gtggcccaca atctttacaa ggtattttta aaaatatcaa agcacaacct 2700 gggaattttg aagcagattc gccaggagat gtggtgcagt atattgctgc acatgataac 2760 cttaccttgc atgatgtgat tgcaaaatca attaataaag accctaaggt agctgaagaa 2820 gatattcata gacgtctgcg tttaggaaat gtaatgattt taacatctca agggacagca 2880 ttcattcatt ctggtcaaga gtatggtcgt acgaagcgtt tacttaaccc tgattacatg 2940 acaaaagttt cagatgacaa attgcctaat aaagcaacac ttattgaagc tgttaaagaa 3000 tacccatatt ttattcatga ttcatatgat tcttcagatg ccattaatca ttttgattgg 3060 gcagcagcca cagataataa caaacaccca atttcaacga aaacacaggc ctatacagca 3120 ggtttaatca cattaaggcg ttcaacagat gctttccgga aattgagcaa agcagaaatt 3180 gatcgtgagg ttagcttgat tacagaggta ggtcaaggtg atattaaaga aaaagatttg 3240 gttattgctt accaaacaat agattctaaa ggcgatattt acgcagtatt tgttaatgct 3300 gatagtaaag ctagaaacgt tttactaggt gaaaaatata aacacctttt aaaagggcaa 3360 gtaattgttg atgctgatca agcggggatt aaaccaatct caactcctag aggtgttcat 3420 tttgaaaaag atagtttgct gattgatcca ttaacagcaa ttgtgattaa agttggcaaa 3480 gttgctccta gccctaagga ggaattgcaa gcagattatc ccaaaacaca atctttcaag 3540 ggatctaaaa cggtagaaaa agtaaataga atagctaata agacctcaat aactcctgta 3600 gtttctaata agaccgattc atatctgaca aatgaagcta atttgccaaa aactggagat 3660 aagtcatcaa aaatactaag tgtagtagga ataagcattc tagcaagtct acttgctcta 3720 3753 ctaggtctct ctttaaagag gaatcgcact taa



<210> 10

<211> 1250

<212> PRT

<213> Streptococcus agalactiae

<400> 10

Met Lys Arg Lys Asp Leu Phe Gly Asp Lys Gln Thr Gln Tyr Thr Ile 

Arg Lys Leu Ser Val Gly Val Ala Ser Val Ala Thr Gly Val Cys Ile 

Phe Leu His Ser Pro Gln Val Phe Ala Glu Glu Val Ser Val Ser Pro 

Ala Thr Thr Ala Ile Ala Lys Ser Asn Ile Asn Gln Val Asp Asn Arg 

Gln Ser Thr Asn Leu Lys Asp Asp Ile Asn Ser Asn Ser Glu Thr Val 70. 

Val Thr Pro Ser Asp Met Pro Asp Thr Lys Gln Leu Val Ser Asp Glu 

Thr Asp Thr Gln Lys Gly Val Thr Glu Pro Asp Lys Ala Thr Ser Leu 

Leu Glu Glu Asn Lys Gly Pro Val Ser Asp Lys Asn Thr Leu Asp Leu 

Lys Val Ala Pro Ser Thr Leu Gln Asn Thr Pro Asp Lys Thr Ser Gln 

Ala Ile Gly Ala Pro Ser Pro Thr Leu Lys Val Ala Asn Gln Ala Pro 

Gln Ile Glu Asn Gly Tyr Phe Arg Leu His Leu Lys Glu Leu Pro Gln 



Gly His Pro Val Glu Ser Thr Gly Leu Trp Ile Trp Gly Asp Val Asp

Gln Pro Ser Ser Asn Trp Pro Asn Gly Ala Ile Pro Met Thr Asn Ala 195 200 205

Lys Lys Asp Asp Tyr Gly Tyr Tyr Val Asp Phe Lys Leu Ser Glu Lys 210 215 220

Gln Arg Lys Gln Ile Ser Phe Leu Ile Asn Asn Lys Ala Gly Thr Asn 225 230 235 240

Leu Ser Gly Asp His His Ile Pro Leu Leu Arg Pro Glu Met Asn Gln
245 250 255

Val Trp Ile Asp Glu Lys Tyr Gly Ile His Thr Tyr Gln Pro Leu Lys
260 265 270

Glu Gly Tyr Val Arg Ile Asn Tyr Leu Ser Ser Ser Gly Asn Tyr Asp
275 280 285

His Leu Ser Ala Trp Leu Phe Lys Asp Val Ala Thr Pro Ser Thr Thr 290 295 300

Trp Pro Asp Gly Ser Asn Phe Val Asn Gln Gly Leu Tyr Gly Arg Tyr
305 310 315 320

Ile Asp Val Pro Leu Lys Thr Asn Ala Lys Glu Ile Gly Phe Leu Ile
325 330 335

Leu Asp Glu Ser Lys Thr Gly Asp Ala Val Lys Val Gln Pro Asn Asp 340 345 350

Tyr Val Phe Arg Asp Leu Ala Asn His Asn Gln Ile Phe Val Lys Asp 355 360 365

Lys Asp Pro Lys Val Tyr Asn Asn Pro Tyr Tyr Ile Asp Gln Val Gln 370 375 380



Leu Lys Asp Ala Gln Gln Thr Asp Leu Thr Ser Ile Gln Ala Ser Phe 385

Thr Thr Leu Asp Gly Val Asp Lys Thr Glu Ile Leu Lys Glu Leu Lys 405 410 415

Val Thr Asp Lys Asn Gln Asn Ala Ile Gln Ile Ser Asp Ile Thr Leu
420 425 430

Asp Thr Ser Lys Ser Leu Leu Ile Ile Lys Gly Asp Phe Asn Pro Lys 435

Gln Gly His Phe Asn Ile Ser Tyr Asn Gly Asn Asn Val Thr Thr Arg
450 455 460

Gln Ser Trp Glu Phe Lys Asp Gln Leu Tyr Ala Tyr Ser Gly Asn Leu 465 470 475 480

Gly Ala Val Leu Asn Gln Asp Gly Ser Lys Val Glu Ala Ser Leu Trp
485 490 . 495

Ser Pro Ser Ala Asp Ser Val Thr Met Ile Ile Tyr Asp Lys Asp Asn 500 505 510

Gln Asn Arg Val Val Ala Thr Thr Pro Leu Val Lys Asn Asn Lys Gly 515 520 525

Val Trp Gln Thr Ile Leu Asp Thr Lys Leu Gly Ile Lys Asn Tyr Thr 530 535 540

Gly Tyr Tyr Tyr Leu Tyr Glu Ile Lys Arg Gly Lys Asp Lys Val Lys 545 550 555 560

Ile Leu Asp Pro Tyr Ala Lys Ser Leu Ala Glu Trp Asp Ser Asn Thr 565 570 575

Val Asn Asp Asp Ile Lys Thr Ala Lys Ala Ala Phe Val Asn Pro Ser 580 585 590



Gln Leu Gly Pro Lys Asn Leu Ser Phe Ala Lys Ile Ala Asn Phe Lys

600

Gly Lys Gln Asp Ala Val Ile Tyr Glu Ala His Val Arg Asp Phe Thr
610 615 620

Ser Asp Gln Ser Leu Asp Gly Lys Leu Lys Asn Gln Leu Gly Thr Phe 625 630 635

Ala Ala Phe Ser Glu Lys Leu Asp Tyr Leu Gln Lys Leu Gly Val Thr
645 650 655

His Ile Gln Leu Leu Pro Val Leu Ser Tyr Phe Tyr Val Asn Glu Met 660 665 670

Asp Lys. Ser Arg Ser Thr Ala Tyr Thr Ser Ser Asp Asn Asn Tyr Asn 675 680 685

Trp Gly Tyr Asp Pro Gln Ser Tyr Phe Ala Leu Ser Gly Met Tyr Ser 690 695 700

Glu Lys Pro Lys Asp Pro Ser Ala Arg Ile Ala Glu Leu Lys Gln Leu 705 710 715 720

Ile His Asp Ile His Lys Arg Gly Met Gly Val Ile Leu Asp Val Val
725 730 735

Tyr Asn His Thr Ala Lys Thr Tyr Leu Phe Glu Asp Ile Glu Pro Asn
740 745 750

Tyr Tyr His Phe Met Asn Glu Asp Gly Ser Pro Arg Glu Ser Phe Gly 755 760 765

Gly Gly Arg Leu Gly Thr Thr His Ala Met Ser Arg Arg Val Leu Val 770 775 780

Asp Ser Ile Lys Tyr Leu Thr Ser Glu Phe Lys Val Asp Gly Phe Arg 785 790 795 795 800



Phe Asp Met Met Gly Asp His Asp Ala Ala Ala Ile Glu Leu Ala Tyr 805 810 815

Lys Glu Ala Lys Ala Ile Asn Pro Asn Met Ile Met Ile Gly Glu Gly 820 825 830

Trp Arg Thr Phe Gln Gly Asp Gln Gly Lys Pro Val Lys Pro Ala Asp 835 840 845

Gln Asp Trp Met Lys Ser Thr Asp Thr Val Gly Val Phe Ser Asp Asp 850 855 860

Ile Arg Asn Ser Leu Lys Ser Gly Phe Pro Asn Glu Gly Thr Pro Ala 865 870 875 880

Phe Ile Thr Gly Gly Pro Gln Ser Leu Gln Gly Ile Phe Lys Asn Ile 885 890 895

Lys Ala Gln Pro Gly Asn Phe Glu Ala Asp Ser Pro Gly Asp Val Val 900 905 910

Gln Tyr Ile Ala Ala His Asp Asn Leu Thr Leu His Asp Val Ile Ala 915 920 925

Lys Ser Ile Asn Lys Asp Pro Lys Val Ala Glu Glu Asp Ile His Arg 930 935 940

Arg Leu Arg Leu Gly Asn Val Met Ile Leu Thr Ser Gln Gly Thr Ala 945 950 955 960

Phe Ile His Ser Gly Gln Glu Tyr Gly Arg Thr Lys Arg Leu Leu Asn 965 970 975

Pro Asp Tyr Met Thr Lys Val Ser Asp Asp Lys Leu Pro Asn Lys Ala 980 985 990

Thr Leu Ile Glu Ala Val Lys Glu Tyr Pro Tyr Phe Ile His Asp Ser 995 1000 1005



Tyr Asp Ser Ser Asp Ala Ile Asn His Phe Asp Trp Ala Ala Ala Thr 1010 1015 .1020

Asp Asn Asn Lys His Pro Ile Ser Thr Lys Thr Gln Ala Tyr Thr Ala 1025 1030 1035 1040

Gly Leu Ile Thr Leu Arg Arg Ser Thr Asp Ala Phe Arg Lys Leu Ser 1045 1050 1055

Lys Ala Glu Ile Asp Arg Glu Val Ser Leu Ile Thr Glu Val Gly Gln
1060 1065 1070

Gly Asp Ile Lys Glu Lys Asp Leu Val Ile Ala Tyr Gln Thr Ile Asp 1075 1080 1085

Ser Lys Gly Asp Ile Tyr Ala Val Phe Val Asn Ala Asp Ser Lys Ala 1090 1095 1100

Arg Asn Val Leu Leu Gly Glu Lys Tyr Lys His Leu Leu Lys Gly Gln 1105 1110 1115 1120

Val Ile Val Asp Ala Asp Gln Ala Gly Ile Lys Pro Ile Ser Thr Pro 1125 1130 1135

Arg Gly Val His Phe Glu Lys Asp Ser Leu Leu Ile Asp Pro Leu Thr

Ala Ile Val Ile Lys Val Gly Lys Val Ala Pro Ser Pro Lys Glu Glu 1155 1160 1165

Leu Gln Ala Asp Tyr Pro Lys Thr Gln Ser Phe Lys Gly Ser Lys Thr 1170 1175 1180

Val Ser Asn Lys Thr Asp Ser Tyr Leu Thr Asn Glu Ala Asn Leu Pro 1205 1210 1215



Lys Thr Gly Asp Lys Ser Ser Lys Ile Leu Ser Val Val Gly Ile Ser 1220 1225 1230

Ile Leu Ala Ser Leu Leu Ala Leu Leu Gly Leu Ser Leu Lys Arg Asn 1235 1240 1245

Arg Thr 1250

<210> 11

<211> 921

<212> DNA

<213> Streptococcus agalactiae

<400> 11

atgaaaaaag tttttttct catggctatg gttgtgagtt tagtaatgat agcagggtgt 60 gataagtcag caaaccccaa acagcctacg caaggcatgt cagttgtaac cagcttttac 120 ccaatgtatg cgatgacaaa agaagtatct ggagacctaa atgatgtgag gatgatccaa 180 tcaggtgcag gcattcattc ctttgaaccg tctgtaaatg atgtggcagc tatttatgac 240 geggatttgt ttgtttacca atcacatacc ttagaagett gggcaaggga tetagaccet 300 aatttaaaaa aatcaaaggt taatgtgttt gaagcgtcaa aacctctgac actagataga 360 gtcaaagggc tagaagatat ggaagtcaca caaggcattg accetgegae actttatgae 420 ccacatacct ggacggatcc cgttttagct ggtgaggaag ctgttaatat cgctaaagag 480 ctaggacatt tggatcctaa acacaaagac agttacacta aaaaggctaa ggctttcaaa 540 aaagaagcag agcaactaac tgaagaatac actcaaaaat ttaaaaaggt gcgctcaaaa 600 . acatttgtga cgcaacaca ggcattttct tatctggcta aacgattcgg cttgaaacaa 660 cttggtatct cgggtatttc tccagagcaa gagccctctc ctcgccaatt gaaagaaatt 720 caagactttg ttaaagaata caacgtcaag actatttttg cagaagacaa cgtcaacccc 780 aaaattgctc atgctattgc gaaatcaaca ggagctaaag taaagacatt aagtccactt 840 gaagctgctc caagcggaaa caagacatat ctagaaaatc ttagagcaaa tttggaagtg 900 ctctatcaac agttgaagta a 921



<210> 12

<211> 306

<212> PRT

<213> Streptococcus agalactiae

<400> 12

Met Lys Lys Val Phe Phe Leu Met Ala Met Val Val Ser Leu Val Met 1 5 10 15

Ile Ala Gly Cys Asp Lys Ser Ala Asn Pro Lys Gln Pro Thr Gln Gly
20 25 30

Met Ser Val Val Thr Ser Phe Tyr Pro Met Tyr Ala Met Thr Lys Glu 35 40 45

Val Ser Gly Asp Leu Asn Asp Val Arg Met Ile Gln Ser Gly Ala Gly
50 55 60

Ile His Ser Phe Glu Pro Ser Val Asn Asp Val Ala Ala Ile Tyr Asp
65 70 75 80

Ala Asp Leu Phe Val Tyr Gln Ser His Thr Leu Glu Ala Trp Ala Arg
85 90 95

Asp Leu Asp Pro Asn Leu Lys Lys Ser Lys Val Asn Val Phe Glu Ala 100 105 110

Ser Lys Pro Leu Thr Leu Asp Arg Val Lys Gly Leu Glu Asp Met Glu 115 120 125

Val Thr Gln Gly Ile Asp Pro Ala Thr Leu Tyr Asp Pro His Thr Trp 130 135 140

Thr Asp Pro Val Leu Ala Gly Glu Glu Ala Val Asn Ile Ala Lys Glu
145 150 155 160

Leu Gly His Leu Asp Pro Lys His Lys Asp Ser Tyr Thr Lys Lys Ala 165 170 175



Lys Ala Phe Lys Lys Glu Ala Glu Gln Leu Thr Glu Glu Tyr Thr Gln 

Lys Phe Lys Lys Val Arg Ser Lys Thr Phe Val Thr Gln His Thr Ala 

Phe Ser Tyr Leu Ala Lys Arg Phe Gly Leu Lys Gln Leu Gly Ile Ser 

Gly Ile Ser Pro Glu Gln Glu Pro Ser Pro Arg Gln Leu Lys Glu Ile 

Gln Asp Phe Val Lys Glu Tyr Asn Val Lys Thr Ile Phe Ala Glu Asp 

Asn Val Asn Pro Lys Ile Ala His Ala Ile Ala Lys Ser Thr Gly Ala 

Lys Val Lys Thr Leu Ser Pro Leu Glu Ala Ala Pro Ser Gly Asn Lys 

Thr Tyr Leu Glu Asn Leu Arg Ala Asn Leu Glu Val Leu Tyr Gln Gln 

Leu Lys 

<210> 13

<211> 657

<212> DNA

<213> Streptococcus agalactiae

<400> 13

ttgttcaata aaataggttt tagaacttgg aaatcaggaa agctttggct ttatatggga 60 gtgctaggat caactattat tttaggatca agtcctgtat ctgctatgga tagtgttgga 120 aatcaaagtc aaggtaatgt tttagagcgt cgccaacgtg atgcggaaaa caaaagtcag 180



657

<210> 14

<211> 218

<212> PRT

<213> Streptococcus agalactiae .

<400> 14

Met Phe Asn Lys Ile Gly Phe Arg Thr Trp Lys Ser Gly Lys Leu Trp

1 5 10 15

ggttttggtt taattttgtt aacaagtcgc tgcggtttga gacgcaatga aaattaa

Leu Tyr Met Gly Val Leu Gly Ser Thr Ile Ile Leu Gly Ser Ser Pro
20 25 30

Val Ser Ala Met Asp Ser Val Gly Asn Gln Ser Gln Gly Asn Val Leu
35 40 45

Glu Arg Arg Gln Arg Asp Ala Glu Asn Lys Ser Gln Gly Asn Val Leu 50 55 60

Glu Arg Arg Gln Arg Asp Ala Glu Asn Lys Ser Gln Gly Asn Val Leu 65 70 75 80

Glu Arg Arg Gln Arg Asp Val Glu Asn Lys Ser Gln Gly Asn Val Leu 85 90 95

Glu Arg Arg Gln Arg Asp Ala Glu Asn Lys Ser Gln Gly Asn Val Leu
100 105 110



Glu Arg Arg Gln Arg Asp Ala Asp Asn Lys Ser Gln Val Gly Gln Leu 115 120 125

Ile Gly Lys Asn Pro Leu Phe Ser Lys Pro Thr Val Ser Arg Glu Asn 130 135 140

Asn His Ser Ser Gln Gly Asp Ser Asn Lys Gln Ser Phe Ser Lys Lys
145 150 155 160

Val Ser Gln Val Thr Asn Val Ala Asn Arg Pro Met Leu Thr Asn Asn 165 170 175

Ser Arg Thr Ile Ser Val Ile Asn Lys Leu Pro Lys Thr Gly Gly Asp 180 185 190

Gln Asn Val Ile Phe Lys Leu Val Gly Phe Gly Leu Ile Leu Leu Thr 195 200 205

Ser Arg Cys Gly Leu Arg Arg Asn Glu Asn 210 215

<210> 15

<211> 1029

<212> DNA

<213> Streptococcus agalactiae

# <400> 15

atgacaaaa aacttatta tgctatata gcactatgca ctatctaac cacttctcáa 60 gctgttttag ctaaagaaa atcacaact gttaccataa aaaacaacta ttcggtctat 120 attaaaaaag aaaaagaga caagccggat aataaaaagc aaatcagcga gacacttaaa 180 gttcctttaa aacccaaaaa agtagttgtt tttgatatgg gagctttgga tactatcaca 240 gctttaggag ctgaaaaatc tgttattggt atcccgaagg ctaaaaatgc tctaaggtta 300 ttgcccaata acgtcaaatc tgttataaa gctaagaga accaagacgt aggaagtctc 360 ttcgaaccaa actttgaagc tattgccga atgcaacctg atgtggttt cctaaggagc 420 cgtatggctt ctgttgataa tattgaaaaa ttaaaagagg ctgcacctaa agcagcatta 480 gtatatgct gagtcgact gagtcgacc aaaaaagta tttgacaaag gagttgctga gcgtgtcaca 540



atgttaggga aaatcttcga ccaaaataa aaggcaaaa cctttaataa agatatcgca 600 caagctgttc ttaaattgca gaaaactatt gagaaaaaag gtaaacctac agctctattt 660 gtaatggcaa acagcggtga acttttaact caatcacctt ctggtcgttt tggttggatt 720 ttctctgtag gtggatttaa agcagtcaat gaaaatgaaa aactaagttc acatggtact 780 cccgtatctt atggaatacat cgctgaaaaa aatcctaact atctttgt tttagatcgt 840 ggagcgacta ttggacaagg agcttcatca aaagaacttt ttaataacga tgttattaaa 900 gcaactgatg ctgtcaaaaa caaacgtgtt catgaggtag atggaaaaga ttggtatatc 960 aattcaggcg gaagccgagt aacactccgt atgattaaag atgtacagaa ctttgttgat 1020 aatcgttaa

<210> 16

<211> 342

<212> PRT

<213> Streptococcus agalactiae

<400> 16

Met Thr Lys Lys Leu Ile Ile Ala Ile Leu Ala Leu Cys Thr Ile Leu 1 5 10 15

Thr Thr Ser Gln Ala Val Leu Ala Lys Glu Lys Ser Gln Thr Val Thr
20 25 30

Ile Lys Asn Asn Tyr Ser Val Tyr Ile Lys Lys Glu Lys Arg Asp Lys
35 40 45

Pro Asp Asn Lys Lys Gln Ile Ser Glu Thr Leu Lys Val Pro Leu Lys
50 55 60

Pro Lys Lys Val Val Val Phe Asp Met Gly Ala Leu Asp Thr Ile Thr
65 70 75 80

Ala Leu Gly Ala Glu Lys Ser Val Ile Gly Ile Pro Lys Ala Lys Asn 85 90 95

Ala Leu Ser Leu Leu Pro Asn Asn Val Lys Ser Val Tyr Lys Ala Lys
100 105 110



Arg Tyr Gln Asp Val Gly Ser Leu Phe Glu Pro Asn Phe Glu Ala Ile 115 120 125

Ala Arg Met Gln Pro Asp Val Val Phe Leu Gly Ala Arg Met Ala Ser 130 135 140

Val Tyr Ala Gly Val Asp Ser Lys Lys Val Phe Asp Lys Gly Val Ala 165 170 175

Glu Arg Val Thr Met Leu Gly Lys Ile Phe Asp Gln Asn Lys Lys Ala 180 185 190

Lys Thr Phe Asn Lys Asp Ile Ala Gln Ala Val Leu Lys Leu Gln Lys
195 200 205

Thr Ile Glu Lys Lys Gly Lys Pro Thr Ala Leu Phe Val Met Ala Asn 210 215 220

Ser Gly Glu Leu Leu Thr Gln Ser Pro Ser Gly Arg Phe Gly Trp Ile 225 230 235 240

Phe Ser Val Gly Gly Phe Lys Ala Val Asn Glu Asn Glu Lys Leu Ser 245 250 255

Ser His Gly Thr Pro Val Ser Tyr Glu Tyr Ile Ala Glu Lys Asn Pro 260 265 270

Asn Tyr Leu Phe Val Leu Asp Arg Gly Ala Thr Ile Gly Gln Gly Ala 275 280 285

Ser Ser Lys Glu Leu Phe Asn Asn Asp Val Ile Lys Ala Thr Asp Ala 290 295 300

Val Lys Asn Lys Arg Val His Glu Val Asp Gly Lys Asp Trp Tyr Ile 305 310 315 315



Asn Ser Gly Gly Ser Arg Val Thr Leu Arg Met Ile Lys Asp Val Gln
325 330 335

Asn Phe Val Asp Asn Arg 340

<210> 17

<211> 2469

<212> DNA

<213> Streptococcus agalactiae

<400> 17

gtgaagaaaa catatggtta tatcggctca gttgctgcta ttttactagc tactcatatt 60 ggaagttacc agcttggtaa gcatcatatg ggtctagcaa caaaggacaa tcagattgcc 120 tatattgatg atagcaaagg taaggtaaaa gcccctaaaa caaacaaaac gatggatcaa 180 atcagtgctg aagaaggcat ctctgctgaa cagatcgtag tcaaaattac tgaccaaggt 240 tatgttacct cacacggtga ccattatcat ttttacaatg ggaaagttcc ttatgatgcg 300 attattagtg aagagttgtt gatgacggat cctaattacc attttaaaca atcagacgtt 360 atcaatgaaa tottagaogg ttaogttatt aaagtcaatg gcaactatta tgtttaccto 420 aagccaggta gtaagcgcaa aaacattcga accaaacaac aaattgctga gcaagtagcc 480 aaaggaacta aagaagctaa agaaaaaggt ttagctcaag tggcccatct cagtaaagaa 540 gaagttgcgg cagtcaatga agcaaaaaga caaggacgct atactacaga cgatggctat 600 atttttagtc cgacagatat cattgatgat ttaggagatg cttatttagt acctcatggt 660 aatcactatc attatattcc taaaaaagat ttgtctccaa gtgagctagc tgctgcacaa 720 gectaetgga gteaaaaaca aggtegaggt getagaeegt etgattaeeg eeegaeaeca 780 gccccaggtc gtaggaaagc cccaattcct gatgtgacgc ctaaccctgg acaaggtcat 840 cagccagata acggtggtta tcatccagcg cctcctaggc caaatgatgc gtcacaaaac 900 aaacaccaaa gagatgagtt taaaggaaaa acctttaagg aacttttaga tcatctacac 960 cgtcttgatt tgaaataccg tcatgtggaa gaagatgggt tgatttttga accgactcaa 1020 gtgatcaaat caaacgcttt tgggtatgtg gtgcctcatg gagatcatta tcatattatc 1080 ccaagaagtc agttatcacc tcttgaaatg gaattagcag atcgatactt agccggccaa 1140 actgatgaca acgactcagg ttcagatcac tcaaaaccat cagataaaga agtgacacat 1200 acctttcttg gtcatcgcat caaagcttac ggaaaaggct tagatggtaa accatatgat 1260 acgagtgatg cttatgtttt tagtaaagaa tccattcatt cagtggataa atcaggagtt 1320 acagctaaac acggagatca tttccactat ataggatttg gagaacttga acaatatgag 1380 ttggatgagg tcgctaactg ggtgaaagca aaaggtcaag ctgatgagct tgttgctgct 1440



ttggatcagg aacaaggcaa agaaaaacca ctctttgaca ctaaaaaagt gagtcgcaaa 1500 gtaacaaaag atggtaaagt gggctatatt atgccaaaag atggcaagga ctatttctat 1560 gctcgttatc aacttgattt gactcagatt gcctttgccg aacaagaact aatgcttaaa 1620 gataagaagc attaccgtta tgacattgtt gatacaggca ttgagccacg acttgctgta 1680 gatgtgtcaa gtctgccgat gcatgctggt aatgctactt acgatactgg aagttcgttt 1740 gttatcccac atattgatca tatccatgtc gttccgtatt catggttgac gcgcaatcag 1800 attgcaacaa tcaagtatgt gatgcaacac cccgaagttc gtccggatgt atggtctaag 1860 ccagggcatg aagagtcagg ttcggtcatt ccaaatgtta cgcctcttga taaacgtgct 1920 ggtatgccaa actggcaaat tatccattct gctgaagaag ttcaaaaagc cctagcagaa 1980 ggtcgttttg cagcaccaga cggctatatt ttcgatccac gagatgtttt ggcaaaagaa 2040 acttttgtat ggaaagatgg ctcctttagc atcccaagag cagatggcag ttcattgaga 2100 accattaata aatccgatct atcccaagct gagtggcaac aagctcaaga gttattggca 2160 aagaaaaatg ctggtgatgc tactgatacg gataaacctg aagaaaagca acaggcagat 2220 aagagcaatg aaaaccaaca gccaagtgaa gccagtaaag aagaaaaaga atcagatgac 2280 tttatagaca gtttaccaga ctatggtcta gatagagcaa ccctagaaga tcatatcaat 2340 caattagcac aaaaagctaa tatcgatcct aagtatctca ttttccaacc agaaggtgtc 2400 caattttata ataaaaatgg tgaattggta acttatgata tcaagacact tcaacaaata 2460 2469 aacccttaa

<210> 18

<211> 822

<212> PRT

<213> Streptococcus agalactiae

<400> 18

Met Lys Lys Thr Tyr Gly Tyr Ile Gly Ser Val Ala Ala Ile Leu Leu 1 5 10 15

Ala Thr His Ile Gly Ser Tyr Gln Leu Gly Lys His His Met Gly Leu
20 25 30

Ala Thr Lys Asp Asn Gln Ile Ala Tyr Ile Asp Asp Ser Lys Gly Lys 35 40 45

Val Lys Ala Pro Lys Thr Asn Lys Thr Met Asp Gln Ile Ser Ala Glu
50 . 55 60



Glu Gly Ile Ser Ala Glu Gln Ile Val Val Lys Ile Thr Asp Gln Gly Tyr Val Thr Ser His Gly Asp His Tyr His Phe Tyr Asn Gly Lys Val Pro Tyr Asp Ala Ile Ile Ser Glu Glu Leu Leu Met Thr Asp Pro Asn Tyr His Phe Lys Gln Ser Asp Val Ile Asn Glu Ile Leu Asp Gly Tyr 1.20 Val Ile Lys Val Asn Gly Asn Tyr Tyr Val Tyr Leu Lys Pro Gly Ser 130 -Lys Arg Lys Asn Ile Arg Thr Lys Gln Gln Ile Ala Glu Gln Val Ala Lys Gly Thr Lys Glu Ala Lys Glu Lys Gly Leu Ala Gln Val Ala His Leu Ser Lys Glu Glu Val Ala Ala Val Asn Glu Ala Lys Arg Gln Gly Arg Tyr Thr Thr Asp Asp Gly Tyr Ile Phe Ser Pro Thr Asp Ile Ile Asp Asp Leu Gly Asp Ala Tyr Leu Val Pro His Gly Asn His Tyr His Tyr Ile Pro Lys Lys Asp Leu Ser Pro Ser Glu Leu Ala Ala Ala Gln 

Ala Tyr Trp Ser Gln Lys Gln Gly Arg Gly Ala Arg Pro Ser Asp Tyr
245 250 255

Arg Pro Thr Pro Ala Pro Gly Arg Lys Ala Pro Ile Pro Asp Val
260 265 270





Thr Pro Asn Pro Gly Gln Gly His Gln Pro Asp Asn Gly Gly Tyr His 

Pro Ala Pro Pro Arg Pro Asn Asp Ala Ser Gln Asn Lys His Gln Arg 

Asp Glu Phe Lys Gly Lys Thr Phe Lys Glu Leu Leu Asp His Leu His 

Arg Leu Asp Leu Lys Tyr Arg His Val Glu Glu Asp Gly Leu Ile Phe 

Glu Pro Thr Gln Val Ile Lys Ser Asn Ala Phe Gly Tyr Val Val Pro 

His Gly Asp His Tyr His Ile Ile Pro Arg Ser Gln Leu Ser Pro Leu 

Glu Met Glu Leu Ala Asp Arg Tyr Leu Ala Gly Gln Thr Asp Asp Asn 

Asp Ser Gly Ser Asp His Ser Lys Pro Ser Asp Lys Glu Val Thr His . 390 

Thr Phe Leu Gly His Arg Ile Lys Ala Tyr Gly Lys Gly Leu Asp Gly 

Lys Pro Tyr Asp Thr Ser Asp Ala Tyr Val Phe Ser Lys Glu Ser Ile 

His Ser Val Asp Lys Ser Gly Val Thr Ala Lys His Gly Asp His Phe 

His Tyr Ile Gly Phe Gly Glu Leu Glu Gln Tyr Glu Leu Asp Glu Val 

Ala Asn Trp Val Lys Ala Lys Gly Gln Ala Asp Glu Leu Val Ala Ala 





_	_	_	_				_		
Leu Asp Gln	Glu Gln	Gly Lys	Glu Lys	Pro Leu	Phe	Asp	Thr	Lys	Lys

Val Ser Arg Lys Val Thr Lys Asp Gly Lys Val Gly Tyr Ile Met Pro 

Lys Asp Gly Lys Asp Tyr Phe Tyr Ala Arg Tyr Gln Leu Asp Leu Thr 

Gln Ile Ala Phe Ala Glu Gln Glu Leu Met Leu Lys Asp Lys Lys His 

Tyr Arg Tyr Asp Ile Val Asp Thr Gly Ile Glu Pro Arg Leu Ala Val 

Asp Val Ser Ser Leu Pro Met His Ala Gly Asn Ala Thr Tyr Asp Thr 

Gly Ser Ser Phe Val Ile Pro His Ile Asp His Ile His Val Val Pro 

Tyr Ser Trp Leu Thr Arg Asn Gln Ile Ala Thr Ile Lys Tyr Val Met 

Gln His Pro Glu Val Arg Pro Asp Val Trp Ser Lys Pro Gly His Glu 

Glu Ser Gly Ser Val Ile Pro Asn Val Thr Pro Leu Asp Lys Arg Ala 

Gly Met Pro Asn Trp Gln Ile Ile His Ser Ala Glu Glu Val Gln Lys 

Ala Leu Ala Glu Gly Arg Phe Ala Ala Pro Asp Gly Tyr Ile Phe Asp 

Pro Arg Asp Val Leu Ala Lys Glu Thr Phe Val Trp Lys Asp Gly Ser 



Phe Ser Ile Pro Arg Ala Asp Gly Ser Ser Leu Arg Thr Ile Asn Lys 690 695 700

Ser Asp Leu Ser Gln Ala Glu Trp Gln Gln Ala Gln Glu Leu Leu Ala 705 710 . 715 720

Lys Lys Asn Ala Gly Asp Ala Thr Asp Thr Asp Lys Pro Glu Glu Lys
725 730 735

Gln Gln Ala Asp Lys Ser Asn Glu Asn Gln Gln Pro Ser Glu Ala Ser 740 745 750

Lys Glu Glu Lys Glu Ser Asp Asp Phe Ile Asp Ser Leu Pro Asp Tyr
755 760 765

Gly Leu Asp Arg Ala Thr Leu Glu Asp His Ile Asn Gln Leu Ala Gln
770 775 780

Lys Ala Asn Ile Asp Pro Lys Tyr Leu Ile Phe Gln Pro Glu Gly Val
785 790 795 800

Gln Phe Tyr Asn Lys Asn Gly Glu Leu Val Thr Tyr Asp Ile Lys Thr 805 810 815

Leu Gln Gln Ile Asn Pro 820

<210> 19

<211> 939

<212> DNA

<213> Streptococcus agalactiae

#### <400> 19

atgatacgcc agtttttaag agaacacttg atttggtata ttttatatat catgatgttt 60 gtcctatttt ttattagttt ctatctatat catttaccaa tgccctattt gtttaattcc 120 ttaggtttaa atgttattgt tttactagga attagtattt ggcaatacag tcgttacagg 180



939

aaaaaatgt tacatctcaa atatttaat agtagtcagg acccctcttt cgaacttcaa 240 ccgagtgatt acgcttattt taatattatt acacaattag aagctagaga agcgcaaaaa 300 gtttctgaaa caattgaaca aaccaatcat gttgcactta tgataaagat gtggtcgcac 360 caaatgaaag ttccattggc agctattca ttaatggccc agacaaatca tctcgatcct 420 aaggaagttg aacaacaatt attgaaattg caacattatc ttgaaacgtt gttagcattt 480 ttgaaattta gacaatatcg tgacgattt cgttttgaag ctgttagcct tagagaagta 540 gtagtagaaa ttataaaatc gtataaggtt atttgctat ccaaaagctt atctatcata 600 attgaaggcg ataatatct ggaaaacagac aaaaagtggt taacttttgc tctttcacag 660 gtgctagata atgccataaa atattctaat cctgagtcaa agataataat aagcatagga 720 gaagagagta ttagaataca agactacggt atcgcatac tcgaagaga tatccctaga 780 ctttttgaag atggctttac gggttacaac ggtcatgagc accaaaaggc aacaggcatg 840 gggttatata tgacaaaaga agtcttatct agtctgaatt tgtccattc ggtggatagc 900

<210> 20

<211> 312

<212> PRT

<213> Streptococcus agalactiae

aaaattaatt atgggactgc tgtttctata cataaataa

<400> 20

Met Ile Arg Gln Phe Leu Arg Glu His Leu Ile Trp Tyr Ile Leu Tyr

1 5 10 15

Ile Met Met Phe Val Leu Phe Phe Ile Ser Phe Tyr Leu Tyr His Leu
20 25 30

Pro Met Pro Tyr Leu Phe Asn Ser Leu Gly Leu Asn Val Ile Val Leu
35 40 45

Leu Gly Ile Ser Ile Trp Gln Tyr Ser Arg Tyr Arg Lys Lys Met Leu 50 55 60

His Leu Lys Tyr Phe Asn Ser Ser Gln Asp Pro Ser Phe Glu Leu Gln
65 70 75 80

Pro Ser Asp Tyr Ala Tyr Phe Asn Ile Ile Thr Gln Leu Glu Ala Arg
85 90 95





Glu Ala Gln Lys Val Ser Glu Thr Ile Glu Gln Thr Asn His Val Ala 

- Leu Met Ile Lys Met Trp Ser His Gln Met Lys Val Pro Leu Ala Ala
- Ile Ser Leu Met Ala Gln Thr Asn His Leu Asp Pro Lys Glu Val Glu
- Gln Gln Leu Leu Lys Leu Gln His Tyr Leu Glu Thr Leu Leu Ala Phe
- Leu Lys Phe Arg Gln Tyr Arg Asp Asp Phe Arg Phe Glu Ala Val Ser
- Leu Arg Glu Val Val Val Glu Ile Ile Lys Ser Tyr Lys Val Ile Cys
- Leu Ser Lys Ser Leu Ser Ile Ile Ile Glu Gly Asp Asn Ile Trp Lys
- Thr Asp Lys Lys Trp Leu Thr Phe Ala Leu Ser Gln Val Leu Asp Asn
- Ala Ile Lys Tyr Ser Asn Pro Glu Ser Lys Ile Ile Ile Ser Ile Gly
- Glu Glu Ser Ile Arg Ile Gln Asp Tyr Gly Ile Gly Ile Leu Glu Glu
- Asp Ile Pro Arg Leu Phe Glu Asp Gly Phe Thr Gly Tyr Asn Gly His
- Glu His Gln Lys Ala Thr Gly Met Gly Leu Tyr Met Thr Lys Glu Val
- Leu Ser Ser Leu Asn Leu Ser Ile Ser Val Asp Ser Lys Ile Asn Tyr



Gly Thr Ala Val Ser Ile His Lys 305 310

<210> 21

<211> 942

<212> DNA

<213> Streptococcus agalactiae

<400> 21

atgacttatc aaaaaacagt tgttttggct ggtgattatt cctacattag acaaattgaa 60 accacattaa aatotototo tototatoat gagaatotot caatttttat tittaatoaa 120 qatatteete aaqaatqqtt tttaqetatg aaagataggg ttggacaaac tggaaatcaa 180 attcaggatg taaagctctt ccatgatcac ttatccccaa aatgggaaaa taaaaagctt 240 aatcatatta attatatqac ctatqctcqt tatttcatac ctcaqtacat ctcaqctqat 300 acaqttttat atcttqactc tqacttagtt gttactacta atttagataa cctctttcaa 360 atttcactag acaatgcata tttagctgca gttccagctc tttttgggct tggatatggg 420 tttaatqctq qaqtaatqqt aattaacaac caacgttggc gacaagaaaa tatgactatt 480 aaattaattq aaaaaaatca aaaggaaatt gagaatgcca acgaagggga tcaaacaatt 540 cttaatcgca tgtttgaaaa tcaggtaatt tatttagatg atacctacaa ttttcaaatt 600 qqttttqata tqqqaqctqc tatcqatqqq cataaattta tttttqacat cccaattacc 660 ccactcccaa aaattattca ctacatttcg ggaatcaaac cttggcaaac attatcaaat 720 atgagactcc gtgaggtatg gtggcactat aatttacttg aatggtcaag tatcatatct 780 agtaaaaaag tatttggttt agaccaccca attaaaacac aaaattatcg tctcaatttc 840 cttattqcta caacttctga ttgtatacca tctatctcag aattagtcac tgcccttcca 900 942 gattgtctat ttcacattgc atgcaccaac agttatgtct ga

<210> 22

<211> 313

<212> PRT

<213> Streptococcus agalactiae

<400> 22

Met Thr Tyr Gln Lys Thr Val Val Leu Ala Gly Asp Tyr Ser Tyr Ile

1 5 10 15



Arg Gln Ile Glu Thr Thr Leu Lys Ser Leu Cys Val Tyr His Glu Asn 

Leu Ser Ile Phe Ile Phe Asn Gln Asp Ile Pro Gln Glu Trp Phe Leu 

Ala Met Lys Asp Arg Val Gly Gln Thr Gly Asn Gln Ile Gln Asp Val 

Lys Leu Phe His Asp His Leu Ser Pro Lys Trp Glu Asn Lys Lys Leu 

Asn His Ile Asn Tyr Met Thr Tyr Ala Arg Tyr Phe Ile Pro Gln Tyr 

Ile Ser Ala Asp Thr Val Leu Tyr Leu Asp Ser Asp Leu Val Val Thr 

Thr Asn Leu Asp Asn Leu Phe Gln Ile Ser Leu Asp Asn Ala Tyr Leu 

Ala Ala Val Pro Ala Leu Phe Gly Leu Gly Tyr Gly Phe Asn Ala Gly 

Val Met Val Ile Asn Asn Gln Arg Trp Arg Gln Glu Asn Met Thr Ile 

Lys Leu Ile Glu Lys Asn Gln Lys Glu Ile Glu Asn Ala Asn Glu Gly 

Asp Gln Thr Ile Leu Asn Arg Met Phe Glu Asn Gln Val Ile Tyr Leu 

Asp Asp Thr Tyr Asn Phe Gln Ile Gly Phe Asp Met Gly Ala Ala Ile 

Asp Gly His Lys Phe Ile Phe Asp Ile Pro Ile Thr Pro Leu Pro Lys 





Ile Ile His Tyr Ile Ser Gly Ile Lys Pro Trp Gln Thr Leu Ser Asn 225 230 235 240

Met Arg Leu Arg Glu Val Trp Trp His Tyr Asn Leu Leu Glu Trp Ser 245 250 255

Ser Ile Ile Ser Ser Lys Lys Val Phe Gly Leu Asp His Pro Ile Lys
260 265 270

Thr Gln Asn Tyr Arg Leu Asn Phe Leu Ile Ala Thr Thr Ser Asp Cys
275 280 285

Ile Pro Ser Ile Ser Glu Leu Val Thr Ala Leu Pro Asp Cys Leu Phe 290 295 300

His Ile Ala Cys Thr Asn Ser Tyr Val 305 310

<210> 23 <211> 1146 <212> DNA

<213> Streptococcus agalactiae

# <400> 23

ggaagttacc agcttggtaa gcatcatatg ggtctagcaa caaaggacaa tcagattgcc 120 tatattgatg atagcaaagg taaggtaaaa gcccctaaaaa caaacaaaac gattggatcaa 180 atagttacct cacacggtga ccattatcat ttttacaatg ggaaagttcc ttatgatgg 300 attattagtg aagagttgtt gatgacggat cctaattacc attttaaaca atcagacgtt 360 atcaatgaaa tcttagacgg ttacgttatt aaagtcaatg gcaactatta tgtttacct 420 aagccaggta gtaagcgcaa aaacattcga accaaacaac aaattgctga gcaagtagcc 480 aaaggaacta aagaagctaa agaaaaagg ttagctcaag tggcccatct cagtaaagaa 540 gaagttgcgg cagtcaatga agcaaaaaga caaggacgct atactacaga cgatggctat 600 attttagtc cgacagatat cattgatgat ttaggagatg cttatttagtc acctaatga accaacaca aaattgctga cagtagcca 540 aattttagtc agcaagatat cattgatgat ttaggagatg cttatttagtc acctcatggt 560 attttagtc attattcc taaaaaagat ttaggagatg cttatttagtc acctcatggt 560 aatcactatc attatattcc taaaaaagat ttggcccaac gtgagctagc tgctgcacaa 720



gcctactgga gtcaaaaca aggtcgaggt gctagaccgt ctgattaccg cccgacacca 780 gccccaggtc gtaggaaagc cccacttcct gatgtgacgc ctaaccctgg acaaggtcat 840 cagccagata acggtggtta tcatccagcg cctcctaggc caaatgatgc gtcacaaaac 900 aaacaccaaa gagatgagtt taaaggaaaa acctttaagg aacttttaga tcaactacac 960 cgtcttgatt tgaaataccg tcatgtggaa gaagatgggt tgattttga accgactcaa 1020 gtgatcaaat caaacgcttt tgggtatgtg gtgcctcatg gagatcatta tcatattatc 1080 ccaagaagtc agttatcacc tcttgaaatg gaattagcag atcgatactt aacccggcca 1140 aactga

<210> 24

<211> 381

<212> PRT

<213> Streptococcus agalactiae

<400> 24

Met Lys Lys Thr Tyr Cys Tyr Ile Gly Ser Val Ala Ala Ile Leu Leu 1 5 10 15

Ala Thr His Ile Gly Ser Tyr Gln Leu Gly Lys His His Met Gly Leu 20 25 30

Ala Thr Lys Asp Asn Gln Ile Ala Tyr Ile Asp Asp Ser Lys Gly Lys
35 40 45

Val Lys Ala Pro Lys Thr Asn Lys Thr Met Asp Gln Ile Ser Ala Glu
50 55 60

Glu Gly Ile Ser Ala Glu Gln Ile Val Val Lys Ile Thr Asp Gln Gly
65 70 75 80

Tyr Val Thr Ser His Gly Asp His Tyr His Phe Tyr Asn Gly Lys Val 85 90 95

Pro Tyr Asp Ala Ile Ile Ser Glu Glu Leu Leu Met Thr Asp Pro Asn 100 105 110





Tyr	His	Phe	Lys	Gln	Ser	Asp	Val	Ile	Asn	Glu	Ile	Leu	Asp	Gly	Tyr
		115					120					125			

- Val Ile Lys Val Asn Gly Asn Tyr Tyr Val Tyr Leu Lys Pro Gly Ser 130 135 140
- Lys Arg Lys Asn Ile Arg Thr Lys Gln Gln Ile Ala Glu Gln Val Ala 145 150 155 160
- Lys Gly Thr Lys Glu Ala Lys Glu Lys Gly Leu Ala Gln Val Ala His

  165 170 175
- Leu Ser Lys Glu Glu Val Ala Ala Val Asn Glu Ala Lys Arg Gln Gly
  180 185 190
- Arg Tyr Thr Thr Asp Asp Gly Tyr Ile Phe Ser Pro Thr Asp Ile Ile 195 200 205
- Asp Asp Leu Gly Asp Ala Tyr Leu Val Pro His Gly Asn His Tyr His 210 215 220
- Tyr Ile Pro Lys Lys Asp Leu Ser Pro Ser Glu Leu Ala Ala Ala Gln 225 230 235 240
- Ala Tyr Trp Ser Gln Lys Gln Gly Arg Gly Ala Arg Pro Ser Asp Tyr
  245 250 255
- Arg Pro Thr Pro Ala Pro Gly Arg Arg Lys Ala Pro Leu Pro Asp Val 260 265 270
- Thr Pro Asn Pro Gly Gln Gly His Gln Pro Asp Asn Gly Gly Tyr His
  275 280 285
- Pro Ala Pro Pro Arg Pro Asn Asp Ala Ser Gln Asn Lys His Gln Arg
  290 295 300
- Asp Glu Phe Lys Gly Lys Thr Phe Lys Glu Leu Leu Asp Gln Leu His 305 310 315 320





Arg Leu Asp Leu Lys Tyr Arg His Val Glu Glu Asp Gly Leu Ile Phe 325 330 335

Glu Pro Thr Gln Val Ile Lys Ser Asn Ala Phe Gly Tyr Val Val Pro 340 345 350

His Gly Asp His Tyr His Ile Ile Pro Arg Ser Gln Leu Ser Pro Leu
355 360 365

Glu Met Glu Leu Ala Asp Arg Tyr Leu Thr Arg Pro Asn 370 375 380

<210> 25

<211> 660

<212> DNA

<213> Streptococcus agalactiae

<400> 25

atggtaaatg atatataga aagaatgtat aaagagaata ttccaaaatc ttaccttaca 60 tccgtcccat tagttattc tcaaaaagga agaacaacct attcgtttag tatgactggt 120 ggtcaacaaa tagatggagt gaaattcaca cagatatatg aggactatat gaaattactc 180 agtcaaggta aggatatcgc agagttatat caaaaatatt ctaaagaaga gttggcaaat 240 ctaggcatta atattatca atccaatgat atacaaagaa ctgaggaaag aacttttgat 300 gaaattatca gttgggtttc caacccttat gcaacaagac caattcaaga aaggcacacct 360 attcaatga agccaacaag atttcacta gaggataaga aacgaattga agaagctgca 420 gctcaaggac taaggcaaat cgaccttatt gattagttg acctatatga tattaatta 480 gacaatacaa gcgtcaatcg ccatattgtg gggttattga ctaatacac ccaagtaaca 540 tactattcc aagaacaatt aaataaggag ttgctgtcaa tggctcaacg tttagataac 600 gtacaacaag cctttattaa attataagt gaagaggaga tacgaaatt tgctctttaa 660

<210> 26

<211> 219

<212> PRT

<213> Streptococcus agalactiae



<400> 26

Met Val Asn Asp Ile Leu Glu Arg Met Tyr Lys Glu Asn Ile Pro Lys

1 10 15

Ser Tyr Leu Thr Ser Val Pro Leu Val Ile Ser Gln Lys Gly Arg Thr
20 25 30

Thr Tyr Ser Phe Ser Met Thr Gly Gly Gln Gln Ile Asp Gly Val Lys
35 40 45

Phe Thr Gln Ile Tyr Glu Asp Tyr Met Lys Leu Leu Ser Gln Gly Lys
50 55 60 .

Asp Ile Ala Glu Leu Tyr Gln Lys Tyr Ser Lys Glu Glu Leu Ala Asn
65 70 75 80

Leu Gly Ile Asn Ile Tyr Gln Ser Asn Asp Ile Glu Arg Thr Glu Glu 85 90 95

Arg Thr Phe Asp Glu Ile Ile Ser Trp Val Ser Asn Pro Tyr Ala Thr
100 105 110

Arg Pro Ile Gln Glu Arg His Thr Ile Gln Leu Glu Pro Thr Arg Phe 115 120 125

Ser Leu Glu Asp Lys Lys Arg Ile Glu Glu Ala Ala Gln Gly Leu 130 135 140

Asp Asn Thr Ser Val Asn Arg His Ile Val Gly Leu Leu Thr Asn Asn 165 170 175

Thr Gln Val Thr Tyr Tyr Phe Gln Glu Gln Leu Asn Lys Glu Leu Leu 180 185 190



44

Ser Met Ala His Ala Leu Asp Asn Val Gln Gln Ala Phe Ile Lys Leu 195 200 205

Leu Ser Glu Glu Glu Ile Arg Lys Phe Ala Leu 210 215

<210> 27

<211> 653

<212> DNA

<213> Streptococcus agalactiae

<400> 27

atgaataaa gaagaaatt atcaaattg aatgtaaaa aacaacttt agcttatgga 60 gctatcactt tagtagccct ttttcatgt attttggctg taacggtcat ctttaaaagt 120 tcacaagtta ctactgaatc tttgtcaaaa gcagataaag ttcgcgtagc caaaaaatca 180 aaaatgacta aggcgacatc taaatcaaaa gtagaagatg taaaacaggc tccaaaacct 240 tctcaggcat ctaatgaagc cccaaaatca agttctcaat ctacagaagc taattctcag 300 caacaagtta ctgcgagtga agaggcggct gtagaacaag cagttgtaac agaaaatacc 360 cctgctacca gtcaggcaa acaaacttat gctgttactg agacaactta caaacctgct 420 caacaccaga caagtggcca agtattgagc aatggaaata ctgcaggggc ggtcggatct 480 gctgctgcag cacaaatgg tgctgcaaca ggagtccctc agtctacttg ggaacatatt 540 attgcccgtg aatcaagtg taatcctaat gttgctaatg cctcaggggc ttcaggact 600 ttccaaaccg tgccaggttg gggttcaaca gctacagttc aggatcaagt taa 653

<210> 28

<211> 234 -

<212> PRT

<213> Streptococcus agalactiae

<400> 28

Met Asn Lys Arg Arg Lys Leu Ser Lys Leu Asn Val Lys Lys Gln His 1 5 10 15

Leu Ala Tyr Gly Ala Ile Thr Leu Val Ala Leu Phe Ser Cys Ile Leu 20 25 30





Ala Val Thr Val Ile Phe Lys Ser Ser Gln Val Thr Thr Glu Ser Leu 35 40 45

45

Ser Lys Ala Asp Lys Val Arg Val Ala Lys Lys Ser Lys Met Thr Lys
50 55 60

Ala Thr Ser Lys Ser Lys Val Glu Asp Val Lys Gln Ala Pro Lys Pro
65 70 75 80

Ser Gln Ala Ser Asn Glu Ala Pro Lys Ser Ser Ser Gln Ser Thr Glu 85 90 95

Ala Asn Ser Gln Gln Gln Val Thr Ala Ser Glu Glu Ala Ala Val Glu
100 105 110

Gln Ala Val Val Thr Glu Asn Thr Pro Ala Thr Ser Gln Ala Gln Gln
115 120 125

Thr Tyr Ala Val Thr Glu Thr Tyr Lys Pro Ala Gln His Gln Thr
130 135 140

Ser Gly Gln Val Leu Ser Asn Gly Asn Thr Ala Gly Ala Val Gly Ser 145 150 155 160

Ala Ala Ala Gln Met Ala Ala Ala Thr Gly Val Pro Gln Ser Thr
165 170 175

Trp Glu His Ile Ile Ala Arg Glu Ser Asn Gly Asn Pro Asn Val Ala 180 185 190

Asn Ala Ser Gly Ala Ser Gly Leu Phe Gln Thr Met Pro Gly Trp Gly
195 200 205

Ser Thr Ala Thr Val Gln Asp Gln Val Asn Ser Ala Ile Lys Ala Tyr 210 215 220

Arg Ala Gln Gly Leu Ser Ala Trp Gly Tyr 225 230



<210> 29

<211> 360

<212> DNA

<213> Streptococcus agalactiae

## <400> 29

atgattgttg gacacggaat tgatttacaa gagatagagg cgattactaa agcatatgag 60 cqtaatcaac gttttgcaga acgcgttttg accgaacaag aattgcttct ttttaaagga 120 atttccaatc ccaagcgtca gatgtctttt ttaacagggc gatgggcagc aaaagaggct 180 tatagcaaag cacttggaac aggaattggg aaagttaatt ttcatgatat cgaaatttta 240 teggatgata aaggagegee tttgattaca aaagaacegt ttaatggaaa atettttgtt 300 tcaatatctc atagtggtaa ttatgcacaa gctagtgtta ttttggagga agaaaaatga 360

<210> 30

<211> 119

<212> PRT

<213> Streptococcus agalactiae

<400> 30

Met Ile Val Gly His Gly Ile Asp Leu Gln Glu Ile Glu Ala Ile Thr 5 10 15 1

Lys Ala Tyr Glu Arg Asn Gln Arg Phe Ala Glu Arg Val Leu Thr Glu 25 30 20

Gln Glu Leu Leu Phe Lys Gly Ile Ser Asn Pro Lys Arg Gln Met 45 40 35

Ser Phe Leu Thr Gly Arg Trp Ala Ala Lys Glu Ala Tyr Ser Lys Ala 60 50 55

Leu Gly Thr Gly Ile Gly Lys Val Asn Phe His Asp Ile Glu Ile Leu 70 75 65



Ser Asp Asp Lys Gly Ala Pro Leu Ile Thr Lys Glu Pro Phe Asn Gly 85 90 95

Lys Ser Phe Val Ser Ile Ser His Ser Gly Asn Tyr Ala Gln Ala Ser 100 105

Val Ile Leu Glu Glu Glu Lys 115

<210> 31

<211> 474

<212> DNA

<213> Streptococcus agalactiae

<400> 31

atgatttttg tcacagtggg gacacatgaa cagcagttca accgtcttat taaagaagtt 60 gatagattaa aagggacagg tgctattgat caagaagtgt tcattcaaac gggttactca 120 gacttcgaac ctcagaattg tcagtggtca aaatttctct catatgatga tatgaactct 180 tacatgaaag aagctgagat tgttatcaca catggcggcc cagcgacgtt tatgtcagtt 240 atttctttag ggaaattacc agttgttgtt cctaggagaa agcagtttgg tgaacatatc 300 aatgatcatc aaatacaatt tttaaaaaaa attgcccacc tgtatccctt ggcttggatt 360 gaagatgtag atggacttgc ggaagcgttg aaaaggaata tagctacaga aaaatatcag 420 ggaaataatg atatgttttg tcataaatta gaaaaaatta taggtgaaat atga

<210> 32

<211> 157

<212> PRT

<213> Streptococcus agalactiae

<400> 32

Met Ile Phe Val Thr Val Gly Thr His Glu Gln Gln Phe Asn Arg Leu 10 15 1

Ile Lys Glu Val Asp Arg Leu Lys Gly Thr Gly Ala Ile Asp Gln Glu



48

Val Phe Ile Gln Thr Gly Tyr Ser Asp Phe Glu Pro Gln Asn Cys Gln 35 40 45

Trp Ser Lys Phe Leu Ser Tyr Asp Asp Met Asn Ser Tyr Met Lys Glu
50 55 60

Ala Glu Ile Val Ile Thr His Gly Gly Pro Ala Thr Phe Met Ser Val 65 70 75 80

Ile Ser Leu Gly Lys Leu Pro Val Val Pro Arg Arg Lys Gln Phe
85 90 95

Gly Glu His Ile Asn Asp His Gln Ile Gln Phe Leu Lys Lys Ile Ala 100 105 110

His Leu Tyr Pro Leu Ala Trp Ile Glu Asp Val Asp Gly Leu Ala Glu 115 120 125

Ala Leu Lys Arg Asn Ile Ala Thr Glu Lys Tyr Gln Gly Asn Asn Asp 130 135 140

Met Phe Cys His Lys Leu Glu Lys Ile Ile Gly Glu Ile 145 150 155

<210> 33

<211> 1203

<212> DNA

<213> Streptococcus agalactiae

### <400> 33

ttggaagaca aattattcaa caaacattt ataggcatta ctatttaaa ctttattgtt 60 tatatggtct attattgtt caccgttatc atagcttta ttgcgactaa agagttaggt 120 gttagcacta gccaagcagg attagcaacg gggatttata ttgtagggac tttgattgct 180 cgtcttatat ttggtaagca attagaagtt ctaggacgta agttagttt acgtggaggg 240 gctattttt acttactaac aactttagct tattttata tgccaagtat cggagtaatg 300 tatttagttc gtttcctaaa tggttttggt tatggcgtcg tgtcaacagc aactaatact 360





attgtaacag cctatatacc agctgataaa agaggtgagg ggattaactt ttacggtcta 420 tcaacaagtt tagccgcagc tattggtcct tttgtaggaa catttatgct agacaacctt 480 catattaact ttaaaatggt tattgtatta tgtagtattt taattgcgat tgtagtgttg 540 qqaqcatttq ttttcccagt caaaaatatt actttaaatc cagaacagtt agctaaatca 600 aaatcatgga ctattgatag tttcattgag aaaaaagcaa tttttatcac aattattgca 660 tttttgatgg gtatctccta tgcttccgtg ttaggtttcc aaaaattata tacaacagaa 720 attaatttga tgacagtagg agcttatttc tttattgttt atgcacttgt catcacttta 780 accagaccat ctatgggaag attaatggac gctaagggag ataagtgggt gctttatcca 840 agttatctgt tcttaacttt gggacttgct ttattaggga gtgctatggg aagtgttacc 900 taccttctat caggtgcttt gattggtttt ggttatggca cctttatgtc ttgtggccaa 960 gcagcatcaa tcaaaggtgt tgaggaacat cgtttcaata cagccatgtc aacttacatg 1020 ataggtettg atttagggtt aggtgetgga cettacattt tgggaettgt taaagatggt 1080 tttcttggag ctggtgtgca atcctttaga gaattattct ggatagcagc gattattcct 1140 gttgtttgtg gtattctata tttcttaaaa tcatctagac aagttgaaac taaaactata 1200 1203 taa

49

<210> 34

<211> 400

<212> PRT

<213> Streptococcus agalactiae

<400> 34

Met Glu Asp Lys Leu Phe Asn Lys His Phe Ile Gly Ile Thr Ile Leu 10 5 15 1

Asn Phe Ile Val Tyr Met Val Tyr Tyr Leu Phe Thr Val Ile Ile Ala 20 25 30

Phe Ile Ala Thr Lys Glu Leu Gly Val Ser Thr Ser Gln Ala Gly Leu 35 40 45

Ala Thr Gly Ile Tyr Ile Val Gly Thr Leu Ile Ala Arg Leu Ile Phe 50 55 60

Gly Lys Gln Leu Glu Val Leu Gly Arg Lys Leu Val Leu Arg Gly Gly 75 80 65 70





Ala Ile Phe Tyr Leu Leu Thr Thr Leu Ala Tyr Phe Tyr Met Pro Ser 85 90 95

Ile Gly Val Met Tyr Leu Val Arg Phe Leu Asn Gly Phe Gly Tyr Gly
100 105 110

Val Val Ser Thr Ala Thr Asn Thr Ile Val Thr Ala Tyr Ile Pro Ala 115 120 125

Asp Lys Arg Gly Glu Gly Ile Asn Phe Tyr Gly Leu Ser Thr Ser Leu 130 135 . 140

Ala Ala Ala Ile Gly Pro Phe Val Gly Thr Phe Met Leu Asp Asn Leu 145 150 155 160

His Ile Asn Phe Lys Met Val Ile Val Leu Cys Ser Ile Leu Ile Ala 165 170 175

Ile Val Val Leu Gly Ala Phe Val Phe Pro Val Lys Asn Ile Thr Leu 180 185 190

Asn Pro Glu Gln Leu Ala Lys Ser Lys Ser Trp Thr Ile Asp Ser Phe 195 200 205

Ile Glu Lys Lys Ala Ile Phe Ile Thr Ile Ile Ala Phe Leu Met Gly
210 215 220

Ile Ser Tyr Ala Ser Val Leu Gly Phe Gln Lys Leu Tyr Thr Thr Glu 225 230 235 240

Ile Asn Leu Met Thr Val Gly Ala Tyr Phe Phe Ile Val Tyr Ala Leu 245 250 255

Val Ile Thr Leu Thr Arg Pro Ser Met Gly Arg Leu Met Asp Ala Lys
260 265 270

Gly Asp Lys Trp Val Leu Tyr Pro Ser Tyr Leu Phe Leu Thr Leu Gly
275 280 285



51

Leu Ala Leu	Leu Gly	Ser	Ala	Met	Gly	Ser	Val	Thr	Tyr	Leu	Leu	Ser
290			295					300				

Gly Ala Leu Ile Gly Phe Gly Tyr Gly Thr Phe Met Ser Cys Gly Gln 305 310 315 320

Ala Ala Ser Ile Lys Gly Val Glu Glu His Arg Phe Asn Thr Ala Met
325 330 335

Ser Thr Tyr Met Ile Gly Leu Asp Leu Gly Leu Gly Ala Gly Pro Tyr 340 345 350

Ile Leu Gly Leu Val Lys Asp Gly Phe Leu Gly Ala Gly Val Gln Ser 355 360 365

Phe Arg Glu Leu Phe Trp Ile Ala Ala Ile Ile Pro Val Val Cys Gly 370 375 380

Ile Leu Tyr Phe Leu Lys Ser Ser Arg Gln Val Glu Thr Lys Thr Ile 385 390 395 400

<210> 35

<211> 393

<212> DNA

<213> Streptococcus agalactiae

## <400> 35

atgaatagtg aacctaaaag tcagtcaaac gaagtaaaaa atagcaagca atcagaagtg 60 aagaaagata aaaaaatgac aaaaaagaa caattagcct atctcaaaga gcatgagcaa 120 gaaatcatag attatgtaaa attacataac aaccaaattg agtccgttca attcgattgg 180 tcaagtgtaa aagtagaaca aagcgggaat ggaactccac aagggggtga ttataatctt 240 tcactgagag gaaagtttaa tcatctacaa aattcaaaat taatagttga tttttattta 300 gctcataaaa atgatatccc aaatatcaaa tcaatgggaa tgctaaataa gccatatata 360

<213> Streptococcus agalactiae

cataaaaatg gtatttggca catttatgaa tag

<400> 36

Met Ile Leu Gly Gly Cys Gln Met Asn Ser Glu Pro Lys Ser Gln Ser

1 5 10 15

Asn Glu Val Lys Asn Ser Lys Gln Ser Glu Val Lys Lys Asp Lys Lys
20 25 30

Met Thr Lys Lys Glu Gln Leu Ala Tyr Leu Lys Glu His Glu Gln Glu 35 40 45

Phe Asp Trp Ser Ser Val Lys Val Glu Gln Ser Gly Asn Gly Thr Pro 65 70 75 80

Gln Gly Gly Asp Tyr Asn Leu Ser Leu Arg Gly Lys Phe Asn His Leu 85 90 95

Gln Asn Ser Lys Leu Ile Val Asp Phe Tyr Leu Ala His Lys Asn Asp 100 105 110

Ile Pro Asn Ile Lys Ser Met Gly Met Leu Asn Lys Pro Tyr Ile His
115 120 125

Lys Asn Gly Ile Trp His Ile Tyr Glu 130 135





<210> 37 <211> 927

<212> DNA

<213> Streptococcus agalactiae

#### <400> 37

atgaaaaaga ttcgattatc aaagtttatt aaaatgattg ttgttatttt gtttttaatt 60 agtgtagcag ctagttttta ttttttccac gttgcccaag ttcgagatga taaatccttt 120 atttcaaatg gtcaacgtaa gcctggaaac tctttatatg cttatgataa atcctttgat 180 aagctattaa agcaaaaaat agaaatgaca aaccaaaata taaagcaagt tgcttggtat 240 gttcctgctg ctaagaaaac tcataagaca gttgttgtcg ttcatggttt tgcgaatagc 300 aaagagaata tgaaggcata tggttggctg tttcataagt taggatacaa tgttcttatg 360 cctgacaaca ttgcacatgg tgaaagtcat gggcagttga taggctatgg ctggaacgac 420 cqcqaqaaca ttatcaaatg gacagaaatg atagtggata agaatccatc aagccaaatt 480 actttatttg gtgtttcaat gggtggagca acagtcatga tggctagtgg tgaaaaatta 540 cctaqtcaqq ttgttaatat cattgaagat tgtggttatt ctagtgtttg ggatgaatta 600 aaatttcagg ctaaagagat gtatggttta ccagccttcc cactcttata tgaagtttca 660 acaatttcta aaatcagagc aggtttttcg tatggacaag caagtagtgt cgaacaattg 720 aaaaagaata atttaccagc cctctttatt catggtgata aggataattt tgttccaaca 780 agtatggttt atgacaacta taaagctaca gcaggtaaga aagagcttta tattgtaaaa 840 ggggcaaaac atgcgaaatc ttttgaaaca gagccagaaa aatatgagaa acgtatctct 900 927 agttttttga aaaaatatga aaaataa

<210> 38

<211> 308

<212> PRT

<213> Streptococcus agalactiae

<400> 38

Met Lys Lys Ile Arg Leu Ser Lys Phe Ile Lys Met Ile Val Val Ile

1 5 10 . 15

Leu Phe Leu Ile Ser Val Ala Ala Ser Phe Tyr Phe Phe His Val Ala
20 25 30

Gln Val Arg Asp Asp Lys Ser Phe Ile Ser Asn Gly Gln Arg Lys Pro





- Gly Asn Ser Leu Tyr Ala Tyr Asp Lys Ser Phe Asp Lys Leu Leu Lys
  50 55 60
- Gln Lys Ile Glu Met Thr Asn Gln Asn Ile Lys Gln Val Ala Trp Tyr
  65 70 75 80
- Val Pro Ala Ala Lys Lys Thr His Lys Thr Val Val Val His Gly
  85 90 95
- Phe Ala Asn Ser Lys Glu Asn Met Lys Ala Tyr Gly Trp Leu Phe His 100 105 110
- Lys Leu Gly Tyr Asn Val Leu Met Pro Asp Asn Ile Ala His Gly Glu 115 120 125
- Ser His Gly Gln Leu Ile Gly Tyr Gly Trp Asn Asp Arg Glu Asn Ile 130 135 140
- Ile Lys Trp Thr Glu Met Ile Val Asp Lys Asn Pro Ser Ser Gln Ile 145 150 155 160
- Thr Leu Phe Gly Val Ser Met Gly Gly Ala Thr Val Met Met Ala Ser 165 170 175
- Gly Glu Lys Leu Pro Ser Gln Val Val Asn Ile Ile Glu Asp Cys Gly 180 185 190
- Tyr Ser Ser Val Trp Asp Glu Leu Lys Phe Gln Ala Lys Glu Met Tyr 195 200 205
- Gly Leu Pro Ala Phe Pro Leu Leu Tyr Glu Val Ser Thr Ile Ser Lys 210 215 220
- Ile Arg Ala Gly Phe Ser Tyr Gly Gln Ala Ser Ser Val Glu Gln Leu 225 230 230 235 235 240
- Lys Lys Asn Asn Leu Pro Ala Leu Phe Ile His Gly Asp Lys Asp Asn 245 250 255





Phe Val Pro Thr Ser Met Val Tyr Asp Asn Tyr Lys Ala Thr Ala Gly
260 265 270

Lys Lys Glu Leu Tyr Ile Val Lys Gly Ala Lys His Ala Lys Ser Phe 275 280 285

Glu Thr Glu Pro Glu Lys Tyr Glu Lys Arg Ile Ser Ser Phe Leu Lys 290 295 300

Lys Tyr Glu Lys 305

<210> 39

<211> 546

<212> DNA

<213> Streptococcus agalactiae

<400> 39

ttgaggagta atatggtaaa gacagcagtt ttaatggcga catacaatgg cgaaaaattt 60 atatctgaac aacttgattc aattcgccaa cagacattaa aaccagatta tgtattattg 120 agggatgatt gttcaacgga tgaaacagtc aatgtcgtca ataactatat cgcaaaacat 180 gagttagaag gctggaaaat tgttaaaaac gacaaaaact taggctggcg tttaaatttt 240 cgtcaattac ttattgatgt gttagcctat gaggttgact atgtctttt tagtgatcaa 300 gatgatattt ggtatcttga taaaaacgaa cgacagtttg ccattatgtc agataaccct 360 caaattgagg ttttgagtgc agacgttgat atcaaaacga tgtctacaga agccagtgtt 420 ccacatttc taacttttc ttctagtgat agaatcagtc agtatcctaa agtatatgat 480 tatcaaaca tccgtccgg atggaccatt gctatgaaga gagattttgc gcaagctatc 540 gcttga



<210> 40

<211> 181

<212> PRT

<213> Streptococcus agalactiae

<400> 40

Met Arg Ser Asn Met Val Lys Thr Ala Val Leu Met Ala Thr Tyr Asn

1 5 10 15

Gly Glu Lys Phe Ile Ser Glu Gln Leu Asp Ser Ile Arg Gln Gln Thr
20 25 30

Leu Lys Pro Asp Tyr Val Leu Leu Arg Asp Asp Cys Ser Thr Asp Glu
35 40 45

Thr Val Asn Val Val Asn Asn Tyr Ile Ala Lys His Glu Leu Glu Gly
50 55 60

Trp Lys Ile Val Lys Asn Asp Lys Asn Leu Gly Trp Arg Leu Asn Phe
65 70 75 80

Arg Gln Leu Leu Ile Asp Val Leu Ala Tyr Glu Val Asp Tyr Val Phe 85 90 95

Phe Ser Asp Gln Asp Asp Ile Trp Tyr Leu Asp Lys Asn Glu Arg Gln
100 105 110

Phe Ala Ile Met Ser Asp Asn Pro Gln Ile Glu Val Leu Ser Ala Asp 115 120 125

Val Asp Ile Lys Thr Met Ser Thr Glu Ala Ser Val Pro His Phe Leu 130 135 140

Thr Phe Ser Ser Ser Asp Arg Ile Ser Gln Tyr Pro Lys Val Tyr Asp 145 150 155 160

Tyr Gln Thr Phe Arg Pro Gly Trp Thr Ile Ala Met Lys Arg Asp Phe 165 170 175



Ala Gln Ala Ile Ala 180

<210> 41

<211> 579

<212> DNA

<213> Streptococcus agalactiae

<400> 41

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<210> 42

<211> 192

<212> PRT

<213> Streptococcus agalactiae

<400> 42

Met Ile His Glu Ile His Asp Cys Gln Phe Ile Glu Lys Gly Ser Tyr

1 5 10 15

Val Tyr Leu Asn Tyr Ile Asn Ala Glu Gly Glu Arg Val Val Ile Ile
20 25 30

Ile Ile Asp Phe Val Arg Ser Val Ser Pro Ile Leu Tyr Arg Leu Phe
35 40 45



58

Met Ile Leu Leu Ala Gln Glu Val Pro His Leu His Asp Tyr Ile Tyr
50 55 60

Asn Ala Arg Asp Asp His Tyr Asp Thr Trp Lys Phe Lys Glu Leu Lys
65 70 75 80

Glu Ser Asn His Pro Val Leu Leu Ala Phe Ser Glu Arg Trp His Asp
85 90 95

Ser Arg Leu Thr Ser Lys Ser Leu Ala Glu Cys Leu Gln Leu Thr Asp 100 105 110

Leu Asp Glu Glu Val Lys Ser Thr Ile Ile Gln Leu Arg Gln Phe Glu 115 120 125

Lys Ser Val Arg Asn Pro Leu Ala His Leu Ile Lys Pro Phe Asp Glu 130 135 140

Gln Glu Leu Tyr Arg Thr Thr Gln Phe Ser Ser Gln Ala Phe Leu Asp 145 150 155 160

Gln Ile Ile Phe Leu Ala Lys Val Ile Gly Val Glu Tyr Asp Thr Val 165 170 175

Asn Phe His Tyr Asp Thr Val Asn Lys Leu Ile Ile Lys Ile Leu Glu 180 185 190

<210> 43

<211> 465

<212> DNA

<213> Streptococcus agalactiae

<400> 43

atggtaaaag tttcaaattt agggtatcca cgtcttggtg aacagcgcga atggaagcaa 60 gcgatcgaag ctttctgggc agggaatctt gaacaaaaag atttagaaaa acaactaaaa 120 caattacgta tcaatcattt aaagaaacaa aaagaggcag gtattgacct tattccagtg 180





<210> 44

<211> 159

<212> PRT

<213> Streptococcus agalactiae

<400> 44

Met Glu Glu Ile Met Val Lys Val Ser Asn Leu Gly Tyr Pro Arg Leu

1 5 10 15

Gly Glu Gln Arg Glu Trp Lys Gln Ala Ile Glu Ala Phe Trp Ala Gly
20 25 30

Asn Leu Glu Gln Lys Asp Leu Glu Lys Gln Leu Lys Gln Leu Arg Ile 35 40 45

Asn His Leu Lys Lys Gln Lys Glu Ala Gly Ile Asp Leu Ile Pro Val

Gly Asp Phe Ser Cys Tyr Asp His Val Leu Asp Leu Ser Phe Gln Phe 65 70 75 80

Asn Val Ile Pro Lys Arg Phe Asp Glu Tyr Glu Arg Asn Leu Asp Leu 85 90 95

Tyr Phe Ala Ile Ala Arg Gly Asp Lys Asp Asn Val Ala Ser Ser Met 100 105 110

Lys Lys Trp Phe Asn Thr Asn Tyr His Tyr Ile Val Pro Glu Trp Glu
115 120 125





Val Glu Thr Lys Pro His Leu Gln Asn Asn Tyr Leu Leu Asp Leu Tyr 135 140 130

Leu Glu Ala Arg Glu Val Val Gly Asp Lys Ala Lys Pro Val Ile 150 155 145

<210> 45

<211> 124

<212> DNA

<213> Streptococcus agalactiae

<400> 45

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<210> 46

<211> 41

<212> PRT

<213> Streptococcus agalactiae

<400> 46

Met Val Leu Leu Leu Leu Met Val Ala Lys Ser Ser Leu Met Val 10 15 1 5

Thr Trp Leu Phe Ile Thr Ile Leu Thr Lys Ile Lys Cys Tyr Gln Ile 25 30 20

Trp Arg Lys Glu Lys Val Ile Lys Leu



<210> 47

<211> 669

<212> DNA

<213> Streptococcus agalactiae

#### <400> 47

atgaacaaa aaatttccgg gatcggcttg gcttcgattg cagtacttag tttagctgca 60
tgtggacatc gtggtgcttc taaatctggt ggtaaatcag atagcttgaa ggttgcaatg 120
gtaacagata ccggtggtgt tgatgataaa tcatttaacc aatctggttg ggaaggtatg 180
caagcttggg gcaagaagaa tggccttaaa aaaggagctg gttttgacta tttccaatcg 240
gcaagtgaat ctgattatgc aactaactta gatacagctg tgtctagtgg ttataaattg 300
attttcggta ttggatttc tctcatgat gctattgata aagcagcaga caataacaaa 360
gatgttaatt acgtcatcgt tgatgatgtt attaaaggga aagataatgt tgcaagtgtt 420
gtctttgcgg ataatgaatc agcttactta gcaggtattg cagccgctaa aactaccaaa 480
acaaaaacag ttggctttgt aggtggtatg gaatctgagg ttattacccg ttttgaaaaa 540
ggttttgaag caggtgtcaa atcagttgat aaatcaatta aaattaaagt tgactatgct 600
ggttcattcg gtgatgctg taagggtaag acaattgcag ccgcacaata tgcttctggc 660
gcagatatt

<210> 48

<211> 223

<212> PRT

<213> Streptococcus agalactiae

### <400> 48

Met Asn Lys Lys Ile Ser Gly Ile Gly Leu Ala Ser Ile Ala Val Leu

1 5 10 15

Ser Leu Ala Ala Cys Gly His Arg Gly Ala Ser Lys Ser Gly Gly Lys
20 25 30

Ser Asp Ser Leu Lys Val Ala Met Val Thr Asp Thr Gly Gly Val Asp
35 40 45

Asp Lys Ser Phe Asn Gln Ser Gly Trp Glu Gly Met Gln Ala Trp Gly
50 55 60





Lys Lys Asn Gly Leu Lys Lys Gly Ala Gly Phe Asp Tyr Phe Gln Ser
65 70 75 80

Ala Ser Glu Ser Asp Tyr Ala Thr Asn Leu Asp Thr Ala Val Ser Ser 85 90 95

Gly Tyr Lys Leu Ile Phe Gly Ile Gly Phe Ser Leu His Asp Ala Ile 100 105 110

Asp Lys Ala Ala Asp Asn Asn Lys Asp Val Asn Tyr Val Ile Val Asp 115 120 125

Asp Val Ile Lys Gly Lys Asp Asn Val Ala Ser Val Val Phe Ala Asp 130 135 140

Asn Glu Ser Ala Tyr Leu Ala Gly Ile Ala Ala Ala Lys Thr Thr Lys 145 150 155 160

Thr Lys Thr Val Gly Phe Val Gly Gly Met Glu Ser Glu Val Ile Thr
165 170 175

Arg Phe Glu Lys Gly Phe Glu Ala Gly Val Lys Ser Val Asp Lys Ser 180 185 190

Ile Lys Ile Lys Val Asp Tyr Ala Gly Ser Phe Gly Asp Ala Ala Lys
195 200 205

Gly Lys Thr Ile Ala Ala Gln Tyr Ala Ser Gly Ala Asp Ile 210 215 220

<210> 49

<211> 609

<212> DNA

<213> Streptococcus agalactiae

<400> 49

atgttacatt ctaaaaaaat acattcctta tcgcttattg ccgttctctc tttagcaaca 60



tatacgagtt tacaaccaaa tcatgtagcg gctgaacaat cacaaaaaac atcaactgtt 120 cttatgagtc aaaaaactat tgaacataag ttaaaagttg cagataaaga agctgctcct 180 ctctacgcta aaatcgacca tatccaacga catattgaag tcaaaaaagc aaaagattta 240 aaagttattg aattgtatat taacaaagat atcaaaccaac tagagaagca aaataaacgt 300 ctactaacta aattctatac ttctattgat aatcaaacat gggatagcac aagtgaagtc 360 aaaaaattga ttgataagac aaccctatcc actaacgaaa aagatagatt aaaaattatat 420 tttgaacaac gtgcttacct tgagacaagg ttgaacgacc gctatcaaaa atttgataac 480 tctattgaa accaaaataa agaactaaaa atattaacgt caaaaataga aaaaatctat 540 caaaaacat gtattacaaa agaggtatta aaaacttact atgctaaaaa aacagtacga 600 gctgactga

<210> 50

<211> 202

<212> PRT

<213> Streptococcus agalactiae

<400> 50

Met Leu His Ser Lys Lys Ile His Ser Leu Ser Leu Ile Ala Val Leu 1 5 10 15

Ser Leu Ala Thr Tyr Thr Ser Leu Gln Pro Asn His Val Ala Ala Glu 20 25 30

Gln Ser Gln Lys Thr Ser Thr Val Leu Met Ser Gln Lys Thr Ile Glu 35 40 45

His Lys Leu Lys Val Ala Asp Lys Glu Ala Ala Pro Leu Tyr Ala Lys
50 55 60

Ile Asp His Ile Gln Arg His Ile Glu Val Lys Lys Ala Lys Asp Leu
65 70 75 80

Lys Val Ile Glu Leu Tyr Ile Asn Lys Asp Ile Asn Gln Leu Glu Lys 85 90 95

Gln Asn Lys Arg Leu Leu Thr Lys Phe Tyr Thr Ser Ile Asp Asn Gln
100 105 110





Thr Trp Asp Ser Thr Ser Glu Val Lys Lys Leu Ile Asp Lys Thr Thr
115 120 125

Leu Ser Thr Asn Glu Lys Asp Arg Leu Lys Leu Tyr Phe Glu Gln Arg 130 135 140

Ala Tyr Leu Glu Thr Arg Leu Asn Asp Arg Tyr Gln Lys Phe Asp Asn 145 150 155 160

Ser Ile Glu Asn Gln Asn Lys Glu Leu Lys Ile Leu Thr Ser Lys Ile
165 170 175

Glu Lys Ile Tyr Gln Lys His Gly Ile Thr Lys Glu Val Leu Lys Thr 180 185 190

Tyr Tyr Ala Lys Lys Thr Val Arg Ala Asp
195 200

<210> 51

<211> 600

<212> DNA

<213> Streptococcus agalactiae

## <400> 51

ctgaattcc aaaaacgcta caatcaaact tggtatccta cttatggtt ttctgatact 60 tatgcattca tggttactaa agagtttgcc agacagaata aaatcaccaa gatctctgat 120 ctcaaaaagt tatcaacaac tatgaaggca ggggttgata gttcatggat gaatcgcgag 180 ggagatggat acactgattt cgctaaaaca tacggttttg aattttcaca tatttaccct 240 atgcaaattg gcttagtcta tgatgcggtt gaaagtaaca aaatgcaatc tgtattaggc 300 tactccactg acggtcgtat ttcgagctat gattagaaa ttttaaggga tgataaaaaa 360 ttctttcctc cttatgaagc ctctatggtt gtcaacaatt ctatcatcaa aaaagatcct 420 aaactaaaaa aattactcca tcgactcgat ggtaaaatca atttaaaaac gatgcaaaac 480 cttaattata tggtagatga taaacttta gaagcttggc gtaatcatgg tcatagctata 600 ttcctgtgtg aaattgttat ccgctcacaa ttccaccaa catacgagcc ggaagcataa 600



<210> 52

<211> 199

<212> PRT

<213> Streptococcus agalactiae

<400> 52

Leu Asn Ser Gln Lys Arg Tyr Asn Gln Thr Trp Tyr Pro Thr Tyr Gly 

Phe Ser Asp Thr Tyr Ala Phe Met Val Thr Lys Glu Phe Ala Arg Gln 

Asn Lys Ile Thr Lys Ile Ser Asp Leu Lys Lys Leu Ser Thr Thr Met 

Lys Ala Gly Val Asp Ser Ser Trp Met Asn Arg Glu Gly Asp Gly Tyr 

Thr Asp Phe Ala Lys Thr Tyr Gly Phe Glu Phe Ser His Ile Tyr Pro 

Met Gln Ile Gly Leu Val Tyr Asp Ala Val Glu Ser Asn Lys Met Gln 

Ser Val Leu Gly Tyr Ser Thr Asp Gly Arg Ile Ser Ser Tyr Asp Leu 

Glu Ile Leu Arg Asp Asp Lys Lys Phe Phe Pro Pro Tyr Glu Ala Ser 

Met Val Val Asn Asn Ser Ile Ile Lys Lys Asp Pro Lys Leu Lys Lys 

Leu Leu His Arg Leu Asp Gly Lys Ile Asn Leu Lys Thr Met Gln Asn 

Leu Asn Tyr Met Val Asp Asp Lys Leu Leu Glu Ala Trp Arg Asn His 





Gly His Ser Cys Phe Leu Cys Glu Ile Val Ile Arg Ser Gln Phe His 180 185 190

Thr Thr Tyr Glu Pro Glu Ala 195

<210> 53

<211> 849

<212> DNA

<213> Streptococcus agalactiae

<400> 53

atqaaaaaat tactttccct aacatgtcta atcatgatgt ctttatgttt agtggcatgt 60 actaaqcaaq caatgtcgtc taagcaagca atgtcgtcta agcaaattaa agataagaat 120 agtaaagaaa aggtgattac tgttgcaact tacagcaaac ctacatctac ctttttagat 180 ttqattaaaq ataatgtaaa agaaaaagga tatactttaa aggttgtcat ggtctctgac 240 tatattcagg ctaacattgc tttagaaaac aaagaacatg atgctaacct tttacaacat 300 gaatttttca tgagtatctt taataaggaa aatgatggtc atctagtgtc aattacacca 360 atttatcatt cattggctgg tttttatggt caacatttga aaaatattgc cgagcttaaa 420 gacggtgcta aggtagcgat tccgtctgat cctgccaata tgactagagc tctgctatta 480 ttgcaagaaa agaaacttat caccttaaag aatacgtcca aaaagaccaa ggctatcgaa 540 gatattatta ctaaccctaa aaaattacga attgaacctg tagcattact taacctcaat 600 caqqcctatt ttqaatatqa ccttqtcttt aatttccctg gatatgtgac aaaaatcaat 660 ctagttccta aaagggatag attattatat gagaaaaaac cagatatccg ttttgcaggt 720 gccttggtag ctcgtgaaga taataaaaat agtgataaaa taaaagtact taaagaagta 780 ctaacaaqta aaqaqattcq tcactatatc actaaggaga ttccaagtga agcagacgtt 840 849 gcgttctag

<210> 54

<211> 282

<212> PRT

<213> Streptococcus agalactiae

<400> 54

Met Lys Lys Leu Leu Ser Leu Thr Cys Leu Ile Met Met Ser Leu Cys

1

5

10





Leu Val Ala Cys Thr Lys Gln Ala Met Ser Ser Lys Gln Ala Met Ser 

- Ser Lys Gln Ile Lys Asp Lys Asn Ser Lys Glu Lys Val Ile Thr Val
- Ala Thr Tyr Ser Lys Pro Thr Ser Thr Phe Leu Asp Leu Ile Lys Asp
- Asn Val Lys Glu Lys Gly Tyr Thr Leu Lys Val Val Met Val Ser Asp
- Tyr Ile Gln Ala Asn Ile Ala Leu Glu Asn Lys Glu His Asp Ala Asn
- Leu Leu Gln His Glu Phe Phe Met Ser Ile Phe Asn Lys Glu Asn Asp
- Gly His Leu Val Ser Ile Thr Pro Ile Tyr His Ser Leu Ala Gly Phe
- Tyr Gly Gln His Leu Lys Asn Ile Ala Glu Leu Lys Asp Gly Ala Lys
- Val Ala Ile Pro Ser Asp Pro Ala Asn Met Thr Arg Ala Leu Leu Leu
- Leu Gln Glu Lys Lys Leu Ile Thr Leu Lys Asn Thr Ser Lys Lys Thr
- Lys Ala Ile Glu Asp Ile Ile Thr Asn Pro Lys Lys Leu Arg Ile Glu
- Pro Val Ala Leu Leu Asn Leu Asn Gln Ala Tyr Phe Glu Tyr Asp Leu
- Val Phe Asn Phe Pro Gly Tyr Val Thr Lys Ile Asn Leu Val Pro Lys



Arg Asp Arg Leu Leu Tyr Glu Lys Lys Pro Asp Ile Arg Phe Ala Gly
225 230 235 240

Ala Leu Val Ala Arg Glu Asp Asn Lys Asn Ser Asp Lys Ile Lys Val 245 250 255

Leu Lys Glu Val Leu Thr Ser Lys Glu Ile Arg His Tyr Ile Thr Lys
260 265 270

Glu Ile Pro Ser Glu Ala Asp Val Ala Phe 275 280

<210> 55

<211> 711

<212> DNA

<213> Streptococcus agalactiae

<400> 55

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<210> 56

<211> 236

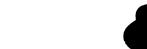
<212> PRT

<213> Streptococcus agalactiae



<400> 56 Leu Leu Ala Lys Glu Thr Thr Met Ser Val Leu Trp Tyr Gln Asn Ser Ala Glu Ala Lys Ala Leu Tyr Leu Gln Gly Tyr Asn Val Ala Lys Met Lys Leu Asp Asp Trp Leu Gln Lys Pro Ser Glu Lys Pro Tyr Ser Ile Ile Leu Asp Leu Asp Glu Thr Val Leu Asp Asn Ser Pro Tyr Gln Ala Lys Asn Ile Lys Asp Gly Ser Ser Phe Thr Pro Glu Ser Trp Asp Lys Trp Val Gln Lys Lys Ser Ala Lys Ala Val Ala Gly Ala Lys Glu Phe Leu Lys Tyr Ala Asn Glu Lys Gly Ile Lys Ile Tyr Tyr Val Ser Asp Arg Thr Asp Ala Gln Val Asp Ala Thr Lys Glu Asn Leu Glu Lys Glu Gly Ile Pro Val Gln Gly Lys Asp His Leu Leu Phe Leu Lys Lys Gly Met Lys Ser Lys Glu Ser Arg Arg Gln Ala Val Gln Lys Asp Thr Asn Leu Ile Met Leu Phe Gly Asp Asn Leu Val Asp Phe Ala Asp Phe Ser Lys Ser Ser Ser Thr Asp Arg Glu Gln Leu Leu Thr Lys Leu Gln Ser Glu Phe Gly Ser Lys Phe Ile Val Phe Pro Asn Pro Met Tyr Gly Ser 

Trp Glu Ser Ala Ile Tyr Gln Gly Lys His Leu Asp Val Gln Lys Gln



Leu Lys Glu Arg Gln Lys Met Leu His Ser Tyr Asp

225

230

235

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<210> 57
<211> 128
<212> DNA
<213> Streptococcus agalactiae
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<400> 57
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ggtgcacaaa tggcttctc aattggtgct agtttgattg cctttgttgg tttagtttct 120
ttgattaa 128

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<210> 58
<211> 42
<212> PRT
<213> Streptococcus agalactiae
<400> 58

Met Asp Asn Lys Gly Asn Asn Ala Asn Val Ile Asp Ala Ile Ala Glu

1 5 10 15
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Gly Ala Ser Thr Gly Ala Gln Met Ala Phe Ser Ile Gly Ala Ser Leu
20 25 30

Ile Ala Phe Val Gly Leu Val Ser Leu Ile 35 40

<210> 59

<211> 573

<212> DNA

<213> Streptococcus agalactiae

<400> 59

atgaaaaga aaaacaaatc ctctaacatt gctataattg caatctttt tgctattatg 60 cttgtcattc atttttgtc atcattatt tttagttttt ggttagtccc tattaaacct 120 actttgatgc atatcccagt tattattgca tctatagcct atggacctcg tattggtgca 180 actctaggcg ccttaatggg ggggatcagc gtagctaaca gcagcattgt tctattacca 240 acgagttacc tcttctcacc ttttgttgaa aatggtaatt tttattcgct aattattgca 300 cttgtaccac gtattctaat cgggattatt ccttattcg tttacaaatt actacacaac 360 cgctttggtt tggctatct aggtgctata ggctctctaa caaacacagt atttgttta 420 tctggaattt ttatctttt ttcaagtact tataatggga atatcaagct aatgctcgct 480 gggattatt catctaattc attagctgag atggtcattg cagctatcat tgtatatcta 540 actgatcctc gtattctca tattaaacat taa

<210> 60

<211> 190

<212> PRT

<213> Streptococcus agalactiae

<400> 60

Met Lys Lys Lys Asn Lys Ser Ser Asn Ile Ala Ile Ile Ala Ile Phe 1 5 10 15

Phe Ala Ile Met Leu Val Ile His Phe Leu Ser Ser Phe Ile Phe Ser





Phe Trp Leu Val Pro Ile Lys Pro Thr Leu Met His Ile Pro Val Ile 35 40 45

Ile Ala Ser Ile Ala Tyr Gly Pro Arg Ile Gly Ala Thr Leu Gly Ala
50 55 60

Leu Met Gly Gly Ile Ser Val Ala Asn Ser Ser Ile Val Leu Pro
65 70 75 80

Thr Ser Tyr Leu Phe Ser Pro Phe Val Glu Asn Gly Asn Phe Tyr Ser

85 90 95

Leu Ile Ile Ala Leu Val Pro Arg Ile Leu Ile Gly Ile Ile Pro Tyr 100 105 110

Phe Val Tyr Lys Leu Leu His Asn Arg Phe Gly Leu Ala Ile Ser Gly
115 120 125

Ala Ile Gly Ser Leu Thr Asn Thr Val Phe Val Leu Ser Gly Ile Phe 130 135 140

Ile Phe Phe Ser Ser Thr Tyr Asn Gly Asn Ile Lys Leu Met Leu Ala .

145 150 155 160

Gly Ile Ile Ser Ser Asn Ser Leu Ala Glu Met Val Ile Ala Ala Ile 165 170 175

Ile Val Tyr Leu Thr Asp Pro Arg Ile Leu Asn Ile Lys His
180 185 190

<210> 61

<211> 251

<212> DNA

<213> Streptococcus agalactiae

<400> 61

ttgaatatga cattacaaga cgaaatcaaa aaacgccgta cttttgccat catctctcac 60





ccggatgctg gtaagacgac tattactgag caattattat attttggtgg tgaaattaga 120 gaagcaggga cagtaaaagg gaaaaaatca ggtacttttg caaagtccga ctggatggat 180 attgaaaagc aacggggtat ctctgttact tcatctgtta tgcaatttga ttacgcgggt 240 aaacgtgtta a

<210> 62

<211> 83

<212> PRT

<213> Streptococcus agalactiae

<400> 62

Met Asn Met Thr Leu Gln Asp Glu Ile Lys Lys Arg Arg Thr Phe Ala

1 5 ' 10 15

Ile Ile Ser His Pro Asp Ala Gly Lys Thr Thr Ile Thr Glu Gln Leu
20 25 30

Leu Tyr Phe Gly Glu Ile Arg Glu Ala Gly Thr Val Lys Gly Lys
35 40 45

Lys Ser Gly Thr Phe Ala Lys Ser Asp Trp Met Asp Ile Glu Lys Gln 50 55 60

Arg Gly Ile Ser Val Thr Ser Ser Val Met Gln Phe Asp Tyr Ala Gly
65 70 75 80

Lys Arg Val

<210> 63

<211> 303

<212> DNA

<213> Streptococcus agalactiae

<400> 63

atggcagata aaaacagaac atttaaactt gtaggtgcag gatcttctag cacacaagaa 60 aaaattgaaa agcctgctct ttcgtttatg caagatgcgt ggcgtcgctt gaaaaaaaac 120





aaattagcag tagtttcact ctatttatta gctcttttac ttacttttc gttagcctca 180 aatttatttg taactcagaa ggatgctaat gggtttgatt cgaaaaaagt aacgacatat 240 cgcaacttac cacctaaatt gagttcaaac cttccttttt ggaatggtag cattaatcca 300 tca

<210> 64

<211> 101

<212> PRT

<213> Streptococcus agalactiae

<400> 64

Met Ala Asp Lys Asn Arg Thr Phe Lys Leu Val Gly Ala Gly Ser Ser

1 5 10 15

Ser Thr Gln Glu Lys Ile Glu Lys Pro Ala Leu Ser Phe Met Gln Asp
20 25 30

Ala Trp Arg Arg Leu Lys Lys Asn Lys Leu Ala Val Val Ser Leu Tyr 35 40 45

Leu Leu Ala Leu Leu Leu Thr Phe Ser Leu Ala Ser Asn Leu Phe Val
50 55 60

Thr Gln Lys Asp Ala Asn Gly Phe Asp Ser Lys Lys Val Thr Thr Tyr
65 70 75 80

Arg Asn Leu Pro Pro Lys Leu Ser Ser Asn Leu Pro Phe Trp Asn Gly
85 90 95

Ser Ile Asn Pro Ser

100



<210> 65

<211> 154 <212> DNA

<213> Streptococcus agalactiae

<400> 65

atgaaaagaa aacagtttat aaaattagga attgcaacct tactaacggt tatttcgctt 60 tacaacaccaa taaacctagc tacaaatcat accacagaaa atattgttac tgctcaagag 120 tataaaacaa agagaatggt actttacctt ttaa 154

<210> 66

<211> 51

<212>, PRT

<213> Streptococcus agalactiae

<400> 66

Met Lys Arg Lys Gln Phe Ile Lys Leu Gly Ile Ala Thr Leu Leu Thr

1 5 10 15

Val Ile Ser Leu Tyr Thr Pro Ile Asn Leu Ala Thr Asn His Thr Thr
20 25 30

Glu Asn Ile Val Thr Ala Gln Glu Tyr Lys Thr Lys Glu Asn Ile Leu
35 40 45

Phe Leu Leu

50

<210> 67

<211> 144

<212> DNA

<213> Streptococcus agalactiae

<400> 67

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<210> 68

<211> 48

<212> PRT

<213> Streptococcus agalactiae

<400> 68

Met Phe Tyr Asn Pro Leu Leu Phe Ile Val Leu Ile Thr Ile Ala Val 1 5 10 15

Phe Phe Leu Ala Lys Lys Trp Gln Leu Pro Thr Phe Thr Phe Ile
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Gly Leu Leu Phe Ile Tyr Asn Gln Gly Leu Trp Glu Gln Leu Ile Asn
35 40 45

<210> 69

<211> 453

<212> DNA

<213> Streptococcus agalactiae

<400> 69

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<210> 70

<211> 150

<212> PRT

<213> Streptococcus agalactiae

<400> 70

Met Val Gln Ile Met Lys Lys His Ile Lys Ser Ile Ile Pro Ile Val 

Leu Ile Gly Met Ile Leu Gly Gly Cys Gln Met Asn Ser Glu His Lys 

Ser Gln Tyr Asn Glu Thr Lys Ser Ser Lys Gln Ser Glu Val Lys Lys 

Asp Lys Lys Met Thr Lys Lys Glu Gln Leu Ala Tyr Leu Lys Glu His 

Glu Gln Glu Ile Ile Asp Phe Val Lys Ser Gln Asn Lys Lys Ile Glu 

Ser Val Gln Ile Asp Trp Asn Asp Val Arg Trp Ser Lys Gly Gly Asn 

Gly Thr Pro Gln Gly Gly Glu Gly Ile Leu Leu Phe Gly Glu Ile 

Asn Asn Asp Ser Glu Ser Ser Trp Arg Val Asp Ile Asp Ile Glu Lys 

Gly Arg Leu Asp Leu Lys Asn Met Tyr Leu Gly Gln Pro Ile Arg Ile 

Gly Gly Lys Leu Phe Glu





<210> 71

<211> 1455

<212> DNA

<213> Streptococcus agalactiae

<400> 71

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qtqqcaqaqc atagaggaca tcatattgat gcattaggga aaaaagattc tacagagaaa 180
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<210> 72

<211> 485

<212> PRT

<213> Streptococcus agalactiae





<400> 72

Met Glu Phe Leu Ala Tyr Asn Ala Phe Thr Ala Ile Gly Val Ser Ile 

Pro His Gly Asn His Phe His Phe Ile His Tyr Lys Asp Met Ser Pro 

Leu Glu Leu Glu Ala Thr Arg Met Val Ala Glu His Arg Gly His His 

Ile Asp Ala Leu Gly Lys Lys Asp Ser Thr Glu Lys Pro Lys His Ile 

Ser His Glu Pro Asn Lys Glu Pro His Thr Glu Glu His His Ala 

Val Thr Pro Lys Asp Gln Arg Lys Gly Lys Pro Asn Ser Gln Ile Val 

Tyr Ser Ala Gln Glu Ile Glu Glu Ala Lys Lys Ala Gly Lys Tyr Thr 

Thr Ser Asp Gly Tyr Ile Phe Asp Ala Lys Asp Ile Lys Lys Asp Thr 

Gly Thr Gly Tyr Val Ile Pro His Met Thr His Glu His Trp Val Pro 

Lys Lys Asp Leu Ser Glu Ser Glu Leu Lys Ala Ala Gln Glu Phe Leu 

Ser Gly Lys Ser Glu Ala Asn Gln Asp Lys Pro Lys Thr Gly Lys Thr 

Ala Gln Glu Ile Tyr Glu Ala Ile Glu Pro Lys Ala Ile Val Lys Pro 





Glu Asp Leu Leu Phe Gly Ile Ala Gln Ala Thr Asp Tyr Lys Asn Gly 

Thr Phe Val Ile Pro His Lys Asp His Tyr His Tyr Val Glu Leu Lys 

Trp Phe Asp Glu Glu Lys Asp Leu Leu Ala Asp Ser Asp Lys Thr Tyr 

Ser Leu Glu Asp Tyr Leu Ala Thr Ala Lys Tyr Tyr Met Met His Pro 

Glu Lys Arg Pro Lys Val Glu Gly Trp Gly Lys Asp Ala Glu Ile Tyr 

Lys Glu Lys Asp Ser Asn Lys Ala Asp Lys Pro Ser Pro Ala Pro Thr 

Asp Asn Lys Ser Thr Ser Asn Ser Ser Asp Lys Asn Leu Ser Ala Ala 

Glu Val Phe Lys Gln Ala Lys Pro Glu Lys Ile Val Pro Leu Asp Lys 

Ile Ala Ala His Met Ala Tyr Ala Val Gly Phe Glu Asp Asp Gln Leu 

Ile Val Pro His His Asp His Tyr His Asn Val Pro Met Ala Trp Phe 

Asp Lys Gly Gly Leu Trp Lys Ala Pro Glu Gly Tyr Thr Leu Gln Gln 

Leu Phe Ser Thr Ile Lys Tyr Tyr Met Glu His Pro Asn Glu Leu Pro 

Lys Glu Lys Gly Trp Gly His Asp Ser Asp His Asn Lys Gly Ser Asn 





Lys Asp Asn Lys Ala Lys Asn Tyr Ala Pro Asp Glu Glu Pro Glu Asp 405 410

81

Ser Gly Lys Val Thr His Asn Tyr Gly Phe Tyr Asp Val Asn Lys Gly 420 425 430

Ser Asp Glu Glu Glu Pro Glu Lys Gln Glu Asp Glu Ser Glu Leu Asp 435 440 445

Glu Tyr Glu Leu Gly Met Ala Gln Asn Ala Lys Lys Tyr Gly Met Asp 455 460 450

Arg Gln Ser Phe Glu Lys Gln Leu Ile Gln Leu Ser Asn Lys Tyr Ser 470 475 480 465

Val Ser Phe Glu Ser 485

<210> 73 <211> 855 <212> DNA <213> Streptococcus agalactiae

## <400> 73

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cacttggttt cagaattagt agatgcaaaa gcagcttcta gtaatgtctt agcttttgaa 780 aaagatggaa aagcttatct tttctcagcc aataacggac gtggcgaagt tgctctttat 840 caattagtaa aataa 855

<210> 74

<211> 284

<212> PRT

<213> Streptococcus agalactiae

<400> 74

Met Arg Lys Arg Phe Ser Leu Leu Asn Phe Ile Val Val Thr Phe Ile

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Phe Phe Phe Ile Leu Phe Pro Leu Phe Lys Ala Lys Asp Cys Gln
20 25 30

Val Val Tyr Ala Ser Phe Gln Gly Asp His Trp Asp Ile Cys Asn Ala 35 40 45

Phe Asp Phe Pro Tyr Leu His Arg Phe Asp Leu Ile Lys Gly Lys Glu 50 55 60

Asn Gln Leu Tyr Phe Ile Gly Cys Thr Ile Ala Asn Ser Lys Ala Tyr
65 70 75 80

Thr Glu Asp Trp Ser Asp Lys Gly Arg Ile Phe Val Ala Arg Phe Asn
85 90 95

Thr Gln Asn His Thr Leu Glu Gly Leu Gln Gln Leu Pro Gln Thr Leu
100 105 110

Leu Lys Asn His Gly Tyr Tyr Ala Ile Gln Asp Glu Gly Tyr Ser Leu 115 120 125

Ile Thr Ser Val Glu Gly Val Leu Lys Leu Thr Tyr Pro Glu Phe Ser 130 135 140





Thr	Thr	Gly	Asp	Trp	Gln	Leu	Glu	Arg	Leu	Phe	Asp	Glu	Glu	Thr	Ser
145					150					155					160

Asp Val Val Lys Val Asp Ile Asn Gln Asp Gly Lys Asp Glu Tyr Val 165 170 175

Ile Ile Gln Gly Phe His Gly Asp Arg Leu Arg Ile Phe Thr Glu Asp 180 185 190

Phe Gly Arg Glu Leu Phe His Tyr Pro Glu Lys Thr Pro Phe Gly His 195 200 205

Ala Ile Trp Ser Gly Arg Leu Leu Asn Gln Thr Cys Phe Val Phe Gly 210 215 220

Trp Arg Ser Glu Lys Ala Glu Leu Arg Leu Phe His Phe Val Asp Gly 230 225 235 240

His Leu Val Ser Glu Leu Val Asp Ala Lys Ala Ala Ser Ser Asn Val 250 245 255

Leu Ala Phe Glu Lys Asp Gly Lys Ala Tyr Leu Phe Ser Ala Asn Asn 260 265 270

Gly Arg Gly Glu Val Ala Leu Tyr Gln Leu Val Lys 275 . 280

<210> 75

<211> 2070

<212> DNA

<213> Streptococcus agalactiae

<400> 75

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tttatcaacc	aatttgtcat	ctcattaaag	gatacaacaa	ttgtatcagc	aatcggactt	2040
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<210> 76

<211> 689

<212> PRT

<213> Streptococcus agalactiae





<400> 76

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Phe Val Phe Gly Gly Lys Val Ser Ala Glu Thr Val Asn Ile Val Ser 

Asp Thr Ala Tyr Ala Pro Phe Glu Phe Lys Asp Ser Asp Gln Thr Tyr 

Lys Gly Ile Asp Val Asp Ile Val Asn Glu Val Ala Lys Arg Ala Gly 

Trp Asn Val Asn Met Thr Tyr Pro Gly Phe Asp Ala Ala Val Asn Ala 

Val Gln Ser Gly Gln Ala Asp Ala Leu Met Ala Gly Thr Thr Val Thr 

Glu Ala Arg Lys Lys Val Phe Asn Phe Ser Asp Thr Tyr Tyr Asp Thr 

Ser Val Ile Leu Tyr Thr Lys Asn Asn Lys Val Thr Asn Tyr Lys 

Gln Leu Lys Gly Lys Val Val Gly Val Lys Asn Gly Thr Ala Ala Gln 

Ser Phe Leu Glu Glu Asn Lys Ser Lys Tyr Gly Tyr Lys Val Lys Thr 

Phe Asp Thr Ser Asp Leu Met Asn Asn Ser Leu Asp Ser Gly Ser Ile 

Tyr Ala Ala Met Asp Asp Gln Pro Val Val Gln Phe Ala Ile Asn Gln 





Gly Lys Ala Tyr Ala Ile Asn Met Glu Gly Glu Ala Val Gly Ser Phe 

Ala Phe Ala Val Lys Lys Gly Ser Gly His Asp Asn Leu Ile Lys Glu 

Phe Asn Thr Ala Phe Ala Gln Met Lys Ser Asp Gly Thr Tyr Asn Asp 

Ile Met Asp Lys Trp Leu Gly Lys Asp Ala Thr Lys Thr Ser Gly Lys 

Ala Thr Gly Asn Ala Asn Glu Lys Ala Thr Pro Val Lys Pro Ser Tyr 

Lys Ile Val Ser Asp Ser Ser Phe Ala Pro Phe Glu Tyr Gln Asn Gly 

Lys Gly Lys Tyr Thr Gly Phe Asp Met Glu Leu Ile Thr Lys Ile Ala 

Lys Gln Gln Gly Phe Lys Leu Asp Ile Ser Asn Pro Gly Phe Asp Ala 

Ala Leu Asn Ala Val Gln Ser Gly Gln Ala Asp Gly Val Ile Ala Gly 

Ala Thr Ile Thr Glu Ala Arg Gln Lys Ile Phe Asp Phe Ser Asp Pro 

Tyr Tyr Thr Ser Ser Val Ile Leu Ala Val Lys Lys Gly Ser Asn Val 

Lys Ser Tyr Gln Asp Leu Lys Gly Lys Thr Val Gly Ala Lys Asn Gly 

Thr Ala Ser Tyr Thr Trp Leu Ser Asp His Ala Asp Lys Tyr Asn Tyr 





His Val Lys Ala Phe Asp Glu Ala Ser Thr Met Tyr Asp Ser Met Asn 

Ser Gly Ser Ile Asp Ala Leu Met Asp Asp Glu Ala Val Leu Ala Tyr 

Ala Ile Asn Gln Gly Arg Lys Phe Glu Thr Pro Ile Lys Gly Glu Lys 

Ser Gly Asp Ile Gly Phe Ala Val Lys Lys Gly Ala Asn Pro Glu Leu 

Ile Lys Met Phe Asn Asn Gly Leu Ala Ser Leu Lys Lys Ser Gly Glu 

Tyr Asp Lys Leu Val Lys Lys Tyr Leu Ser Thr Ala Ser Thr Ser Ser 

Asn Asp Lys Ala Ala Lys Pro Val Asp Glu Ser Thr Ile Leu Gly Leu 

Ile Ser Asn Asn Tyr Lys Gln Leu Leu Ser Gly Ile Gly Thr Thr Leu 

· Ser Leu Thr Leu Ile Ser Phe Ala Ile Ala Met Val Ile Gly Ile Ile 

Phe Gly Met Met Ser Val Ser Pro Ser Asn Thr Leu Arg Thr Ile Ser 

Met Ile Phe Val Asp Ile Val Arg Gly Ile Pro Leu Met Ile Val Ala 

Ala Phe Ile Phe Trp Gly Ile Pro Asn Leu Ile Glu Ser Ile Thr Gly 

His Gln Ser Pro Ile Asn Asp Phe Val Ala Ala Thr Ile Ala Leu Ser 



88

Leu Asn Gly Gly Ala Tyr Ile Ala Glu Ile Val Arg Gly Gly Ile Glu 610 620

Ala Val Pro Ser Gly Gln Met Glu Ala Ser Arg Ser Leu Gly Ile Ser 625 630 635 640

Tyr Gly Lys Thr Met Gln Lys Val Ile Leu Pro Gln Ala Val Arg Leu 645 650 655

Met Leu Pro Asn Phe Ile Asn Gln Phe Val Ile Ser Leu Lys Asp Thr
660 665 670

Thr Ile Val Ser Ala Ile Gly Leu Val Glu Leu Phe Gln Thr Gly Lys
675 680 685

Ser

<210> 77 <211> 149

<212> DNA

<213> Streptococcus agalactiae

<400> 77

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<210> 78

<211> 49

<212> PRT

<213> Streptococcus agalactiae

<400> 78

Met Glu Gly Leu Leu Ile Ala Leu Ile Pro Met Phe Ala Trp Gly Ser

10

1 5

15





Ile Gly Phe Val Ser Asn Lys Ile Gly Gly Arg Pro Asn Gln Gln Thr
20 25 30

Phe Gly Met Thr Leu Gly Ala Leu Leu Phe Ala Ile Ile Val Cys Leu 35 40 45

Phe

<210> 79

<211> 963

<212> DNA

<213> Streptococcus agalactiae

<400> 79

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Ala Leu Asp Ser Tyr Leu Asp Asn Ile Asn Phe Asp Tyr Thr Val Thr 

Asp Ile Ser Asp Glu Tyr Lys Met Leu Gln Ile Glu Gly Arg Tyr Ser 

Gly Glu Ile Ala Gln Ser Phe Tyr Glu Tyr Asp Ile Ser Thr Leu Asn 

Phe Arg Thr Leu Arg Ile Glu Met Asp Phe Ile Lys Gly Glu Glu Arg 

Leu Ser Trp Arg Arg Phe Gly Phe Ser Gly Glu Phe Gly Tyr Gln Phe 

Phe Leu Pro Ser Ser Ile Phe Ala Thr Phe Val Ser Asp Val Cys Glu 





Gly Ile Ala Glu Cys Gly Asp Glu Leu Asp Arg Tyr Leu Arg Phe Glu 

Val Gly Gln Pro Ile Thr Asp Ile Tyr Gln Glu Glu Tyr Ser Leu 

Tyr Glu Ile Gly Tyr Ser Trp Asn Leu Asp Phe Thr Lys Glu Glu Phe 

Arg Gly Arg Asp Ser Leu Leu Glu His Ile Arg Ser Ala Thr Val Lys 

Ser Val Gly Phe Ser Thr Lys Glu Lys Leu Ala Ser Gly Thr Pro Val 

Leu Phe Asp Asp Gln Ile Val Gly Lys Ile Phe Trp Ile Ala Asp Glu 

Lys His Ser Ser Glu Asn Tyr Leu Gly Leu Met Ile Val Asn Gln Thr 

Tyr Ala His Ser Gly Val Thr Phe Val Thr Glu Asp Gly Gln Ile Leu 

Lys Thr Gln Ser Ser Pro Tyr Cys Ile Pro Glu Ser Trp Asn Lys Glu 

<210> 81

<211> 702

<212> DNA

<213> Streptococcus agalactiae

## <400> 81

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<210> 82

<211> 233

<212> PRT

<213> Streptococcus agalactiae

<400> 82

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Val Leu Arg Gly Ala Ser Tyr Arg Phe Tyr Ser Gly Lys Ile Thr Gly 20 25 30

Val Leu Gly Arg Asn Gly Ala Gly Lys Thr Thr Leu Phe Asn Ile Leu 35 40 45

Tyr Gly Asp Leu Ala Ala Asp Asn Gly Thr Ile Cys Leu Leu Lys Asp 55 50

Asn His Glu Tyr Pro Leu Thr Asp Lys Asp Ile Gly Ile Val Tyr Ser 70 75 80 65

Glu Asn Tyr Leu Pro Glu Phe Leu Thr Gly Tyr Glu Phe Val Lys Phe 85 90 95

Tyr Met Asp Leu His Pro Ser Asp Asp Leu Met Thr Ile Asp Asp Tyr 105 110 100



Leu Asp Phe Met Glu Ile Gly Gln Thr Glu Arg His Arg Ile Ile Lys
115 120 125

Gly Tyr Ser Asp Gly Met Lys Ser Lys Leu Ser Leu Ile Cys Leu Met 130 135 140

Ile Ser Lys Pro Lys Val Ile Leu Leu Asp Glu Pro Leu Thr Ala Val
145 150 155 160

Asp Val Val Ser Ser Ile Ala Ile Lys Arg Leu Leu Glu Leu Ser 165 170 175

Glu Asp His Ile Ile Ile Leu Ser Thr His Ile Met Ala Leu Ala Glu 180 185 190

Asp Leu Cys Asp Ile Val Ala Val Leu Asp Lys Gly Lys Leu Gln Thr
195 200 205

Leu Asp Ile Asp Arg Lys His Glu Gln Phe Glu Glu Arg Leu Leu Gln
210 220

Val Leu Lys Gly Asp Glu Tyr Asp Lys 225 230

<210> 83

<211> 774

<212> DNA

<213> Streptococcus agalactiae

#### <400> 83

ttgtttatga gatatacaaa tggaaatttt gaagcetttg caagacetcg aaaacetgaa 60 ggtgtggata aaaaateege ttatattgtt ggttetggtt tageaggatt agetgeeget 120 gtetttttaa taegtgaegg teaaatggat ggteaaegta tteatatttt tgaagaaeta 180 eetetttetg gaggateaet tgaeggtgte aaaegaeetg atateggttt tgtaaegegt 240 ggtggtegtg aaatggaaaa teaettegaa tgtatgtggg atatgtaeeg tteeateeee 300 teetetegaag tteeagatge tteetateta gatgaatttt attggettga eaaggatgat 360



cccaattcat ctaactgtcg cctcattcat aaacagggga atcgcttaga atctgatggt 420 gattttacac tcggaacaca ttccaaagag ttagttaagc tagtcatgga gactgaagag 480 tctttaggtg ctaagacgat tgaagaagtt ttttcaaaag aatttttga aagtaatttt 540 tggacttatt gggctactat gtttgccttt gagaaatggc attcagcgat tgaaatgcgt 600 cgatatgcta tgcgctttat ccatcatatt ggtggtctgc ctgatttcac ttcattaaaa 660 tttaataaat ataatcaata tgattctatg gtgaaaccaa tcatcagtta tttagagtct 720 cataatgtag atgttcaatt tgatagcaag gtaactaata tctccgtaga cttt 774

<210> 84

<211> 258

<212> PRT

<213> Streptococcus agalactiae

<400> 84

Met Phe Met Arg Tyr Thr Asn Gly Asn Phe Glu Ala Phe Ala Arg Pro 1 5 10 15

Arg Lys Pro Glu Gly Val Asp Lys Lys Ser Ala Tyr Ile Val Gly Ser
20 25 30

Gly Leu Ala Gly Leu Ala Ala Ala Val Phe Leu Ile Arg Asp Gly Gln
35 40 45

Met Asp Gly Gln Arg Ile His Ile Phe Glu Glu Leu Pro Leu Ser Gly
50 55 60

Gly Ser Leu Asp Gly Val Lys Arg Pro Asp Ile Gly Phe Val Thr Arg
65 70 75 80

Gly Gly Arg Glu Met Glu Asn His Phe Glu Cys Met Trp Asp Met Tyr

85 90 95

Arg Ser Ile Pro Ser Leu Glu Val Pro Asp Ala Ser Tyr Leu Asp Glu 100 105 110

Phe Tyr Trp Leu Asp Lys Asp Asp Pro Asn Ser Ser Asn Cys Arg Leu 115 120 125



Ile His Lys Gln Gly Asn Arg Leu Glu Ser Asp Gly Asp Phe Thr Leu 130 135 140

Gly Thr His Ser Lys Glu Leu Val Lys Leu Val Met Glu Thr Glu Glu 145 150 155 160

Ser Leu Gly Ala Lys Thr Ile Glu Glu Val Phe Ser Lys Glu Phe Phe 165 170 175

Glu Ser Asn Phe Trp Thr Tyr Trp Ala Thr Met Phe Ala Phe Glu Lys
180 185 190

Trp His Ser Ala Ile Glu Met Arg Arg Tyr Ala Met Arg Phe Ile His
195 200 205

His Ile Gly Gly Leu Pro Asp Phe Thr Ser Leu Lys Phe Asn Lys Tyr 210 215 220

Asn Gln Tyr Asp Ser Met Val Lys Pro Ile Ile Ser Tyr Leu Glu Ser 225 230 235 240

His Asn Val Asp Val Gln Phe Asp Ser Lys Val Thr Asn Ile Ser Val
245 250 255

Asp Phe

<210> 85

<211> 903

<212> DNA

<213> Streptococcus agalactiae

## <400> 85

ttgttggctt ctttatttat cgtccgtttg tcaaaatcgc tttcgctaag gaggagcaat 60 atgaaaaaat tacttagatg gcttcctcct gtacttttca ttattatcct tataggaatg 120 actatcttag gtaagtccta tatcaataaa gtaacagctc acaaaataaa actctataac 180





tctcgaatga ctcctactat tttaatttca ggatccagtg ctactcaaga acgatttaac 240 agcatgttag cacageteaa eeaaatggga gaaaaacata gegttttaaa gttaaetgte 300 aaaaaagaca atagcattat ctacaatgga caaattagcg gcaatgacca caaaccctac 360 attqtcattq gatttgaaaa taatgaagat ggttatagta acatcaaaaa acaaacaaaa 420 tggctacaga ttgctatgaa tgatcttcag aagaaatata aatttaaacg ttttaacgct 480 atcqqtcatt caaatqqtqq cttatcatqq actattttcc tagaaqatta ttacqactct 540 gatgaatttg atatgaaatc attgttaaca atgggaacac cttttaactt tgaagaaagt 600 aacacctcaa atcatactca aatgcttaaa gatttaatca gtaataaagg aaatattcca 660 tcaagtctca tggtatacaa tttggcagga actaattcat atgatggtga taaaattgtt 720 ccatttgcta gtgtggagac tggtaaatat attttccaag aaaccgctaa acactatacc 780 caactaacag taactggtaa taatgctaca cattctgact tgcctgataa tcctgaagtt 840 atccaatatg tcgcagaaaa aattcttaaa aatgagaaag gtaaattacc aaaacctcac 900 903

<210> 86

taa

<211> 300

<212> PRT

<213> Streptococcus agalactiae

<400> 86

Met Leu Ala Ser Leu Phe Ile Val Arg Leu Ser Lys Ser Leu Ser Leu 5 10 15 1

Arg Arg Ser Asn Met Lys Lys Leu Leu Arg Trp Leu Pro Pro Val Leu 30 20 25

Phe Ile Ile Leu Ile Gly Met Thr Ile Leu Gly Lys Ser Tyr Ile 45 40 35

Asn Lys Val Thr Ala His Lys Ile Lys Leu Tyr Asn Ser Arg Met Thr 60 50 55

Pro Thr Ile Leu Ile Ser Gly Ser Ser Ala Thr Gln Glu Arg Phe Asn 70 75 80 65

Ser Met Leu Ala Gln Leu Asn Gln Met Gly Glu Lys His Ser Val Leu 95 85 90



Lys Leu Thr Val Lys Lys Asp Asn Ser Ile Ile Tyr Asn Gly Gln Ile 100 105 110

Ser Gly Asn Asp His Lys Pro Tyr Ile Val Ile Gly Phe Glu Asn Asn 115 120 125

Glu Asp Gly Tyr Ser Asn Ile Lys Lys Gln Thr Lys Trp Leu Gln Ile 130 135 140

Ile Gly His Ser Asn Gly Gly Leu Ser Trp Thr Ile Phe Leu Glu Asp 165 170 175

Tyr Tyr Asp Ser Asp Glu Phe Asp Met Lys Ser Leu Leu Thr Met Gly
180 185 190

Thr Pro Phe Asn Phe Glu Glu Ser Asn Thr Ser Asn His Thr Gln Met
195 200 205

Leu Lys Asp Leu Ile Ser Asn Lys Gly Asn Ile Pro Ser Ser Leu Met 210 215 220

Val Tyr Asn Leu Ala Gly Thr Asn Ser Tyr Asp Gly Asp Lys Ile Val 225 230 235 240

Pro Phe Ala Ser Val Glu Thr Gly Lys Tyr Ile Phe Gln Glu Thr Ala 245 250 255

Lys His Tyr Thr Gln Leu Thr Val Thr Gly Asn Asn Ala Thr His Ser
260 265 270

Asp Leu Pro Asp Asn Pro Glu Val Ile Gln Tyr Val Ala Glu Lys Ile 275 280 285

Leu Lys Asn Glu Lys Gly Lys Leu Pro Lys Pro His
290 295 300



<210> 87 <211> 912 <212> DNA <213> Streptococcus agalactiae

<400> 87

ttqaaattag gtattacaac attcggagag acaacaatcc ttgaagaaac aaaccaaagc 60 tattcacatc ctgagaggat tcgccaatta gttgctgaga ttgaactagc tgatcaagtt 120 qqtttaqatg tatatggtat tggagagcac catcgtgaag attttgcggt ctctgcaccc 180 qaaattatcc tagcagcagg agcggttaga actaataata tccgtttatc tagtgcagta 240 acquittetet ettecaatga teetattege gtetateage aatttteaac gattgaegea 300 ctttcaaatq qtaqaqcaqa aattatggca gggcgtggtt cctttattga gtcttttcca 360 ttgtttggat acgatttagc ggattatgat gatttattta atgaaaaaat ggatatgttg 420 ttagcaatta actcagcgac aaatctcgat tggaaaggtc atttgacaca aacagttaat 480 qaqcqaccaa tttatccaag agcattacaa agacagttat caatatgggt ggcaacagga 540 qqaaatqttq attctacaat tcgtattgca gaacaaggtt tgccaattgt ttatgcaact 600 attqqtqqqa atcccaaaqc ctttcqtcaa ttgqtccata tttataaaga agttggtaag 660 tccgtaatgg acacaaacca ggaacaacta aaagttgctg ctcactcttg gggatggatt 720 gaagaggata atcaaaccgc tattgaccgt tattttttcc ctacgaaaca gaccgtcgat 780 aatattgcta agggacgccc tcattggtct gaaatgacta aagagcagta tttacgttca 840 ataggtccag aaggtgctat ttttgtagga aatcctgaag tggttgcaca taaaattata 900 912 ggactttggt ga

<210> 88

<211> 303

<212> PRT

<213> Streptococcus agalactiae

<400> 88

Met Lys Leu Gly Ile Thr Thr Phe Gly Glu Thr Thr Ile Leu Glu Glu

1 5 10 15

Thr Asn Gln Ser Tyr Ser His Pro Glu Arg Ile Arg Gln Leu Val Ala

20





Glu Ile Glu Leu Ala Asp Gln Val Gly Leu Asp Val Tyr Gly Ile Gly
35 40 45

Glu His His Arg Glu Asp Phe Ala Val Ser Ala Pro Glu Ile Ile Leu 50 55 60

Ala Ala Gly Ala Val Arg Thr Asn Asn Ile Arg Leu Ser Ser Ala Val 65 70 75 80

Thr Ile Leu Ser Ser Asn Asp Pro Ile Arg Val Tyr Gln Gln Phe Ser 85 90 95

Thr Ile Asp Ala Leu Ser Asn Gly Arg Ala Glu Ile Met Ala Gly Arg 100 105 110

Gly Ser Phe Ile Glu Ser Phe Pro Leu Phe Gly Tyr Asp Leu Ala Asp 115 120 125

Tyr Asp Asp Leu Phe Asn Glu Lys Met Asp Met Leu Leu Ala Ile Asn 130 135 140

Glu Arg Pro Ile Tyr Pro Arg Ala Leu Gln Arg Gln Leu Ser Ile Trp 165 170 175

Val Ala Thr Gly Gly Asn Val Asp Ser Thr Ile Arg Ile Ala Glu Gln 180 185 190

Gly Leu Pro Ile Val Tyr Ala Thr Ile Gly Gly Asn Pro Lys Ala Phe 195 200 205

Arg Gln Leu Val His Ile Tyr Lys Glu Val Gly Lys Ser Val Met Asp 210 215 220

Thr Asn Gln Glu Gln Leu Lys Val Ala Ala His Ser Trp Gly Trp Ile
225 230 235 240



Glu Glu Asp Asn Gln Thr Ala Ile Asp Arg Tyr Phe Phe Pro Thr Lys
245 250 255

Gln Thr Val Asp Asn Ile Ala Lys Gly Arg Pro His Trp Ser Glu Met
260 265 270

Thr Lys Glu Gln Tyr Leu Arg Ser Ile Gly Pro Glu Gly Ala Ile Phe 275 280 285

Val Gly Asn Pro Glu Val Val Ala His Lys Ile Ile Gly Leu Trp 290 295 300

<210> 89

<211> 693

<212> DNA

<213> Streptococcus agalactiae

<400> 89

atgatagagt ggattcaaac acatttacca aatgtatatc aaatgggttg ggaaggtgct 60 tacggctggc agacagctat tgtacaaacc ctttatatga ctttttggtc gttccttatt 120 ggaggtttaa tggaggttta ttccttgtt taactagtcc tagaggaggtt 180 attgctaata aattagtatt tggagttta gataaagttg tttctgttt tagagctctg 240 cccttcatta ttcttcttgc tttgattgcg ccagtaactc gcgtaattgt aggaacaaca 300 cttggttcac cagcagcttt ggtacctctt tctttggcag ttttcccatt ttttgctcgt 360 caagttcaag ttgttttagc tgaacttgat ggtggagtta ttgaggctgc acaagcctca 420 ggtggaacac tttgggatat tattgtagtt tatcttcgtg aaggtctacc agatttaatt 480 cgagtatcaa cggttacttt gattcttta gtaggtgaaa cagctatggc tggcgctatt 540 ggtgcaggag gattgggtc tggtgctatt actaaaggat ataactatc tcgtgatgat 600 attactttag tagcgcctt gagtcataaa taattaattt tctttatcca attttaggt 660 gattttttaa cacgtcgctt gagtcataaa taa

<210> 90

<211> 230

<212> PRT

<213> Streptococcus agalactiae



101

<400> 90

Met Ile Glu Trp Ile Gln Thr His Leu Pro Asn Val Tyr Gln Met Gly

1 5 10 15

Trp Glu Gly Ala Tyr Gly Trp Gln Thr Ala Ile Val Gln Thr Leu Tyr
20 25 30

Met Thr Phe Trp Ser Phe Leu Ile Gly Gly Leu Met Gly Leu Leu Gly
35 40 45

Gly Leu Phe Leu Val Leu Thr Ser Pro Arg Gly Val Ile Ala Asn Lys
50 55 60

Leu Val Phe Gly Val Leu Asp Lys Val Val Ser Val Phe Arg Ala Leu 65 70 75 80

Pro Phe Ile Ile Leu Leu Ala Leu Ile Ala Pro Val Thr Arg Val Ile 85 90 95

Val Gly Thr Thr Leu Gly Ser Pro Ala Ala Leu Val Pro Leu Ser Leu
100 105 110

Ala Val Phe Pro Phe Phe Ala Arg Gln Val Gln Val Leu Ala Glu 115 120 125

Leu Asp Gly Gly Val Ile Glu Ala Ala Gln Ala Ser Gly Gly Thr Leu 130 135 140

Trp Asp Ile Ile Val Val Tyr Leu Arg Glu Gly Leu Pro Asp Leu Ile 145 150 155 160

Arg Val Ser Thr Val Thr Leu Ile Ser Leu Val Gly Glu Thr Ala Met 165 170 175

Ala Gly Ala Ile Gly Ala Gly Gly Leu Gly Ser Val Ala Ile Thr Lys 180 185 190



102

Gly Tyr Asn Tyr Ser Arg Asp Asp Ile Thr Leu Val Ala Thr Ile Leu 195 200 205

Ile Leu Leu Leu Ile Phe Phe Ile Gln Phe Leu Gly Asp Phe Leu Thr
210 215 220

Arg Arg Leu Ser His Lys 225 230

<210> 91

<211> 759

<212> DNA

<213> Streptococcus agalactiae

<400> 91

ttggcagtta gtttcatga agtatttggt tgggattctg cttttttat tatgattatc 60 aatattccat tgctccttct ttgctacttt ggcttaggta aacaaacctt tttaaaaact 120 gtctatggtt cttggatttt tcctgttttt attaagttaa cacaaagtgt accaactttg 180 acccacaact cactcctcgc agcacttttt ggaggtgtta ttgtaggatg tggtttgggg 240 attgttttt ggagcgactc ttcaactggt ggaacgggga ttatcattca attcttagga 300 aaatatactc ctataagcct tggacaaggg gttatattga ttggtggact tgttacaatt 360 gttggtttcc tagcttttga cagtgatacg gttatgttt ctattattgg gttgataact 420 attagttata ttattaatgc tatccaaact ggatttacaa ccttaagcac tgtcttaatc 480 gtttctcaag agcaccaaaa aattaagaca tatatcaata ctgtcgcaga tagaggagta 540 acagaaattc ccgttaaagg gggatattct ggaactaatc aaatcatgct tatgacaact 600 attgctggtt atgagttgc taaattacaa gaggcaatag cagaaattga cgaaacagcc 660 ttcataacag tagatgaaga cattcttatg ccaatgtaa ccaatgta tcaaaaaa 720 catggacgtc ttgatgaaga cattcttatg ccaatgtaa



103

<210> 92

<211> 252

<212> PRT

<213> Streptococcus agalactiae

<400> 92

Met Ala Val Ser Phe His Glu Val Phe Gly Trp Asp Ser Ala Phe Phe 1 5 10 15

Ile Met Ile Ile Asn Ile Pro Leu Leu Leu Cys Tyr Phe Gly Leu
20 25 30

Gly Lys Gln Thr Phe Leu Lys Thr Val Tyr Gly Ser Trp Ile Phe Pro 35 40 45

Val Phe Ile Lys Leu Thr Gln Ser Val Pro Thr Leu Thr His Asn Ser
50 55 60

Leu Leu Ala Ala Leu Phe Gly Gly Val Ile Val Gly Cys Gly Leu Gly 65 70 75 80

Ile Val Phe Trp Ser Asp Ser Ser Thr Gly Gly Thr Gly Ile Ile
85 90 95

Gln Phe Leu Gly Lys Tyr Thr Pro Ile Ser Leu Gly Gln Gly Val Ile 100 105 110

Leu Ile Asp Gly Leu Val Thr Ile Val Gly Phe Leu Ala Phe Asp Ser 115 120 125

Asp Thr Val Met Phe Ser Ile Ile Gly Leu Ile Thr Ile Ser Tyr Ile 130 135 140

Ile Asn Ala Ile Gln Thr Gly Phe Thr Thr Leu Ser Thr Val Leu Ile 145 150 155 160

Val Ser Gln Glu His Gln Lys Ile Lys Thr Tyr Ile Asn Thr Val Ala 165 170 175



Asp Arg Gly Val Thr Glu Ile Pro Val Lys Gly Gly Tyr Ser Gly Thr 180 185 190

Asn Gln Ile Met Leu Met Thr Thr Ile Ala Gly Tyr Glu Phe Ala Lys

195 \_200 205

Leu Gln Glu Ala Ile Ala Glu Ile Asp Glu Thr Ala Phe Ile Thr Val 210 215 220

Thr Pro Thr Ser Gln Ala Ser Gly Arg Gly Phe Ser Leu Gln Lys Asn 225 230 235 240

His Gly Arg Leu Asp Glu Asp Ile Leu Met Pro Met
245 250

<210> 93 <211> 549

<212> DNA

<213> Streptococcus agalactiae

# <400> 93

atgaaagaaa aacagtcgaa aaggcttatt tatactac tgattgtcc cattatcttt 60 ataagtgttt ttacatacag tattagccag cettctaaac tacttccacc aaaagaatta 120 gttattctaa gtccaaatag tcaagccatt ttaacaggaa cgattccagc ttttgaggaa 180 aaatacggta taaaagttaa gcttattcaa ggtgggacag ggcaactaat agatagatta 240 agtaaggagg gtaagcagtt gaaggcggat atttcttg gaggaaatta tacgcaattt 300 gaaagtcata aggcattgtt tgagtcttac gtatcaaaga atgttcatac tgttattcca 360 gactatatcc atccgagtga tacggcgaca ccttatacta taaatgggag tgtcttgatt 420 gtaaaataacg aattagctaa gggacttacc atcaagagtt atgaagattt attacagcct 480 tccttaaaag gtaaaattgc ctttgcagat cctctagagt cgacctgcaa gcatgcaagc 540 ttggcgtaa



105

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<	っ	1	О	>	9	4

<211> 182

<212> PRT

<213> Streptococcus agalactiae

#### <400> 94

Met Lys Glu Lys Gln Ser Lys Arg Leu Ile Tyr Ile Leu Leu Ile Val

Pro Ile Ile Phe Ile Ser Val Phe Thr Tyr Ser Ile Ser Gln Pro Ser
20 25 30

Lys Leu Leu Pro Pro Lys Glu Leu Val Ile Leu Ser Pro Asn Ser Gln 35 40 45

Ala Ile Leu Thr Gly Thr Ile Pro Ala Phe Glu Glu Lys Tyr Gly Ile
50 55 60

Lys Val Lys Leu Ile Gln Gly Gly Thr Gly Gln Leu Ile Asp Arg Leu 65 70 75 80

Ser Lys Glu Gly Lys Gln Leu Lys Ala Asp Ile Phe Phe Gly Gly Asn 85 90 95

Tyr Thr Gln Phe Glu Ser His Lys Ala Leu Phe Glu Ser Tyr Val Ser 100 105 110

Lys Asn Val His Thr Val Ile Pro Asp Tyr Ile His Pro Ser Asp Thr 115 120 125

Ala Thr Pro Tyr Thr Ile Asn Gly Ser Val Leu Ile Val Asn Asn Glu 130 135 140

Leu Ala Lys Gly Leu Thr Ile Lys Ser Tyr Glu Asp Leu Leu Gln Pro 145 150 155 160

Ser Leu Lys Gly Lys Ile Ala Phe Ala Asp Pro Leu Glu Ser Thr Cys 165 170 175



106

Lys His Ala Ser Leu Ala 180

<210> 95

<211> 368

<212> DNA

<213> Streptococcus agalactiae

<400> 95

cctcctatca aatgatgaca aacgtgagag gtacatggaa caaatgctct ttaaaattga 60 aaatgcaacc tggcagcgtg tggtaagagc actttatcgt aaatacaata aggaatttt 120 tacatatcca gccgccaaaa caaaccacca cgcttttgaa tcaggattgg catatcacac 180 ggcaacaatg gttcgtttgg cagatagtat cggagatatc tatccagaac ttaataaaag 240 tttgatgttt gctggtatta tgctacatga tttagccaag gtcatagagt tatcgggtcc 300 tgataataca gaatatacta ttcgaggtaa tcttatcggt catattcac ttattgatga 360 ggaattaa

<210> 96

<211> 122

<212> PRT

<213> Streptococcus agalactiae

<400> 96

Leu Leu Ser Asn Asp Asp Lys Arg Glu Arg Tyr Met Glu Gln Met Leu

1 5 10 15

Phe Lys Ile Glu Asn Ala Thr Trp Gln Arg Val Val Arg Ala Leu Tyr
20 25 30

Arg Lys Tyr Asn Lys Glu Phe Phe Thr Tyr Pro Ala Ala Lys Thr Asn
35 40 45

His His Ala Phe Glu Ser Gly Leu Ala Tyr His Thr Ala Thr Met Val 50 55 60



Arg Leu Ala Asp Ser Ile Gly Asp Ile Tyr Pro Glu Leu Asn Lys Ser
65 70 75 80

Leu Met Phe Ala Gly Ile Met Leu His Asp Leu Ala Lys Val Ile Glu 85 90 95

Leu Ser Gly Pro Asp Asn Thr Glu Tyr Thr Ile Arg Gly Asn Leu Ile
100 105 110

Gly His Ile Ser Leu Ile Asp Glu Glu Leu 115 120

<210> 97

<211> 753

<212> DNA

<213> Streptococcus agalactiae

<400> 97

atgaaaaaa ataaattat ccgattcagt ttagttggt ttetacttgc gatactatgc 60 tttagtcttt ttgctttatt gaagcctaac agtcaacaat catcatcta aaagttgagg 120 aatgaggata taaaaaagac atcctctaa aaaagaaata agaaattacg attaccagct 180 gtatcatcaa aagattggaa cttgatttg gtcaatcgtg accataaaca tgaagaatta 240 agtccagatg tggtgcctgt tgaaaatat tatttggata aacgtattac gaagcaagct 300 actcagttt tagaggctgc tagagcaatt gattcacgag aacatttaat ttcgggttat 360 cgtagtgttg cctatcagga gaagttgttc aattctatg ttaccaaga gatgactagt 420 aaccctaatt tgacgagggg acaagcagaa aagttggtaa aaacttactc tcagcctgca 480 ggtgctagtg aacaccagac tggattagcg atggatatga gtactgtaga ttctttgaat 540 gagagcgatc ctagagtagt cagtcagttg aacaagcaga gtactgaag gttatgaaga ttggcattac 660 cgctatgtt gggtagagtc tgcaaaatat atggtcaaca atcatttaac attagaagaa 720 tacataactt tattaaagga gaataaccaa tgg



<210> 98

<211> 250

<212> PRT

<213> Streptococcus agalactiae

<400> 98

Met Lys Lys Asn Lys Ile Ile Arg Phe Ser Leu Val Gly Val Leu Leu

1 5 10 15

Ala Ile Leu Cys Phe Ser Leu Phe Ala Leu Leu Lys Pro Asn Ser Gln
20 25 30

Gln Ser Ser Gln Lys Leu Arg Asn Glu Asp Ile Lys Lys Thr Ser
35 40 45

Ser Gln Lys Arg Asn Lys Lys Leu Arg Leu Pro Ala Val Ser Ser Lys
50 55 60

Asp Trp Asn Leu Ile Leu Val Asn Arg Asp His Lys His Glu Glu Leu 65 70 75 80

Ser Pro Asp Val Val Pro Val Glu Asn Ile Tyr Leu Asp Lys Arg Ile 85 90 95

Thr Lys Gln Ala Thr Gln Phe Leu Glu Ala Ala Arg Ala Ile Asp Ser 100 105 110

Arg Glu His Leu Ile Ser Gly Tyr Arg Ser Val Ala Tyr Gln Glu Lys
115 120 125

Leu Phe Asn Ser Tyr Val Thr Gln Glu Met Thr Ser Asn Pro Asn Leu 130 135 140

Thr Arg Gly Gln Ala Glu Lys Leu Val Lys Thr Tyr Ser Gln Pro Ala 145 150 155 160

Gly Ala Ser Glu His Gln Thr Gly Leu Ala Met Asp Met Ser Thr Val 165 170 175



Asp Ser Leu Asn Glu Ser Asp Pro Arg Val Val Ser Gln Leu Lys Lys 180 185 190

Ile Ala Pro Gln Tyr Gly Phe Val Leu Arg Phe Pro Asp Gly Lys Thr
195 200 205

Ala Glu Thr Gly Val Gly Tyr Glu Asp Trp His Tyr Arg Tyr Val Gly
210 215 220

Val Glu Ser Ala Lys Tyr Met Val Lys His His Leu Thr Leu Glu Glu 225 230 235 240

Tyr Ile Thr Leu Leu Lys Glu Asn Asn Gln 245 250

<210> 99

<211> 351

<212> DNA

<213> Streptococcus agalactiae

<400> 99

ctgttatgtg gatttette ateaatteet gtgtetaatt eegggggta tggtataata 60 acagttatga aaaataaaaa aatettattt gggactggee ttgetggtg gggtttaetg 120 geagetgetg gttataeeet aaetaaaaaa gtaacagatt ataaaegtea geaaateaet 180 cagacettaa gagaaetttt tagteagatg ggtgatatte aggtatttta ttttaatgaa 240 tttgaatetg atattaaaat gaceagtggt ggtettgtet tggaagatgg cagaattte 300 gaatteattt ategteaagg tgttettgat tatgtggagg tgageaaatg a 351

<210> 100

<211> 116

<212> PRT

<213> Streptococcus agalactiae

<400> 100

Leu Leu Cys Gly Phe Leu Pro Ser Ile Pro Val Ser Asn Ser Gly Gly

1 5 10 15



110

Tyr Gly Ile Ile Thr Val Met Lys Asn Lys Lys Ile Leu Phe Gly Thr
20 25 30

Gly Leu Ala Gly Val Gly Leu Leu Ala Ala Gly Tyr Thr Leu Thr
35 40 45

Lys Lys Val Thr Asp Tyr Lys Arg Gln Gln Ile Thr Gln Thr Leu Arg
50 55 60

Glu Leu Phe Ser Gln Met Gly Asp Ile Gln Val Phe Tyr Phe Asn Glu
65 70 75 80

Phe Glu Ser Asp Ile Lys Met Thr Ser Gly Gly Leu Val Leu Glu Asp
85 90 95

Gly Arg Ile Phe Glu Phe Ile Tyr Arg Gln Gly Val Leu Asp Tyr Val
100 105 110

Glu Val Ser Lys 115

<210> 101

<211> 310

<212> DNA

<213> Streptococcus agalactiae

<400> 101

atgtatcaaa ctcagacaaa taaggaaaaa tttgtttat ttttgaaatt atttatcca 60 gtattgattt atcaatttgc taattttca gctactttta ttgattcggt tatgactgga 120 cagtatagtc agctacattt ggcaggtgtg tcaactgcta gtaatttatg gactccgttt 180 ttcgcttat tagtaggtat gattcagca ttagtaccag tagttggtca acatttgggt 240 agaggaaata aagaacaaat tcgcacagaa tttcatcaat ttctatattt aggtttgata 300 ctgtccttaa



<210> 102

<211> 103

<212> PRT

<213> Streptococcus agalactiae

<400> 102

Met Tyr Gln Thr Gln Thr Asn Lys Glu Lys Phe Val Leu Phe Leu Lys

1 5 10 15

Leu Phe Ile Pro Val Leu Ile Tyr Gln Phe Ala Asn Phe Ser Ala Thr
20 25 30

Phe Ile Asp Ser Val Met Thr Gly Gln Tyr Ser Gln Leu His Leu Ala 35 40 45

Gly Val Ser Thr Ala Ser Asn Leu Trp Thr Pro Phe Phe Ala Leu Leu 50 55 60

Val Gly Met Ile Ser Ala Leu Val Pro Val Val Gly Gln His Leu Gly 65 70 75 80

Arg Gly Asn Lys Glu Gln Ile Arg Thr Glu Phe His Gln Phe Leu Tyr 85 90 95

Leu Gly Leu Ile Leu Ser Leu 100

<210> 103

<211> 1098

<212> DNA

<213> Streptococcus agalactiae

<400> 103

ctgctctttt tagctaactt ttctaattta tggtataatt gtatggattg tttagctaga 60 atggagaaga tgatgcaaga tgttttcatt ataggaagta gagggttgcc agctcgttac 120 ggtggttttg aaacttttgt ttcagaattg attaatcatc aaaaaagttc cgacataaaa 180

taccatqttq catqccttaq tqataaaqaa catcatactc attttaactt tqctgacqct 240 gattqtttta ctataaatcc tccccaatta gggccagcac gtgtgattgc ttatgatatt 300 atggccatta attatgccct tgacttggtt aagacacatg atttaaaaga gcctattttt 360 tatattttag gaaatacaat tggtgccttt atttggcatt ttgccaataa aatacataaa 420 qtcggtggct tattgtatgt taatccggat ggtttagagt ggaagcgatc aaagtggtct 480 cqtcccacac aqcqttattt aaaatacqcc gaaaaatqta tqactaaaaa tqcaqaccta 540 attatttctg ataatattgg tattgaaaat tacattcaat ctacctactc taatgtgaag 600 acaaggttca ttgcttacgg tacagagatt aattctagga aattatcgtc agatgatcca 660 cqtqtcaaac agttgtttaa aaaatggaat attaagtcta agggttacta tctaatcgtt 720 ggtcgatttg tccctgaaaa caattatgaa acggctatta gggagttcat ggcttcagat 780 actaagcgtg atttagttat tatctgtaac catcaaaata acccctactt tgaaaagttg 840 tccttaaaqa caaaccttca acaagataaa agagttaagt ttgtaggtac gctctatgaa 900 aaaqatctqc tqgattatqt tcqtcaacaa qcctttqctt atattcatqq qcatqaaqtt 960 ggcggtacta atccaggact gcttgaggct ttagctaata ctgatttgaa tcttgttcta 1020 qatqttgatt tcaacaaatc agtagcaggt ctctcaagtt tttactggac taaaaaagag 1080 ggggatttag ctaagctt 1098

<210> 104

<211> 366

<212> PRT

<213> Streptococcus agalactiae

<400> 104

Met Leu Phe Leu Ala Asn Phe Ser Asn Leu Trp Tyr Asn Cys Met Asp

1 5 10 15

Cys Leu Ala Arg Met Glu Lys Met Met Gln Asp Val Phe Ile Ile Gly
20 25 30

Ser Arg Gly Leu Pro Ala Arg Tyr Gly Gly Phe Glu Thr Phe Val Ser 35 40 45

Glu Leu Ile Asn His Gln Lys Ser Ser Asp Ile Lys Tyr His Val Ala 50 55 60

Cys Leu Ser Asp Lys Glu His His Thr His Phe Asn Phe Ala Asp Ala 65 70 75 80



113 Asp Cys Phe Thr Ile Asn Pro Pro Gln Leu Gly Pro Ala Arg Val Ile 95 85 90 Ala Tyr Asp Ile Met Ala Ile Asn Tyr Ala Leu Asp Leu Val Lys Thr 100 105 110 His Asp Leu Lys Glu Pro Ile Phe Tyr Ile Leu Gly Asn Thr Ile Gly 120 115 Ala Phe Ile Trp His Phe Ala Asn Lys Ile His Lys Val Gly Gly Leu 130 135 140 Leu Tyr Val Asn Pro Asp Gly Leu Glu Trp Lys Arg Ser Lys Trp Ser 150 155 160 145 Arg Pro Thr Gln Arg Tyr Leu Lys Tyr Ala Glu Lys Cys Met Thr Lys 165 170 175 Asn Ala Asp Leu Ile Ile Ser Asp Asn Ile Gly Ile Glu Asn Tyr Ile 190 185 180 Gln Ser Thr Tyr Ser Asn Val Lys Thr Arg Phe Ile Ala Tyr Gly Thr

Gln Ser Thr Tyr Ser Asn Val Lys Thr Arg Phe Ile Ala Tyr Gly Thr 195 200 205

Glu Ile Asn Ser Arg Lys Leu Ser Ser Asp Asp Pro Arg Val Lys Gln 210 215 220

Leu Phe Lys Lys Trp Asn Ile Lys Ser Lys Gly Tyr Tyr Leu Ile Val 225 230 235 240

Gly Arg Phe Val Pro Glu Asn Asn Tyr Glu Thr Ala Ile Arg Glu Phe 245 250 255

Met Ala Ser Asp Thr Lys Arg Asp Leu Val Ile Ile Cys Asn His Gln 260 265 270

Asn Asn Pro Tyr Phe Glu Lys Leu Ser Leu Lys Thr Asn Leu Gln Gln 275 280 285





Asp	Lys	Arg	Val	Lys	Phe	Val	Gly	Thr	Leu	Tyr	G <sup>·</sup> lu	Lys	Asp	Leu	Leu
	290					295					300				

Asp Tyr Val Arg Gln Gln Ala Phe Ala Tyr Ile His Gly His Glu Val 305 310 315 320

Gly Gly Thr Asn Pro Gly Leu Leu Glu Ala Leu Ala Asn Thr Asp Leu
325 330 335

Asn Leu Val Leu Asp Val Asp Phe Asn Lys Ser Val Ala Gly Leu Ser
340 345 350

Ser Phe Tyr Trp Thr Lys Lys Glu Gly Asp Leu Ala Lys Leu 355 360 365

<210> 105

<211> 546

<212> DNA

<213> Streptococcus agalactiae

<400> 105

ttgaggagta atatggtaaa gacagcagtt ttaatggcga catacaatgg cgaaaaattt 60 atatctgaac aacttgattc aattcgccaa cagacattaa aaccagatta tgtattattg 120 agggatgatt gttcaacgga tgaaacagtc aatgtcgtca ataactatat cgcaaaacat 180 gagttagaag gctggaaaat tgttaaaaac gacaaaaact taggctggcg tttaaatttt 240 cgtcaattac ttattgatgt gttagcctat gaggttgact atgtctttt tagtgatcaa 300 gatgatattt ggtatcttga taaaaacgaa cgacagtttg ccattatgtc agataaccct 360 caaattgagg ttttgagtgc agacgttgat atcaaaacga tgtctacaga agccagtgtt 420 ccacatttc taacttttc ttctagtgat agaatcagtc agtatcctaa agtatatgat 480 tatcaaacat tccgtcccgg atggaccatt gctatgaaga gagattttgc gcaagctatc 540 gcttga



<210> 106

<211> 181

<212> PRT

<213> Streptococcus agalactiae

<400> 106

Met Arg Ser Asn Met Val Lys Thr Ala Val Leu Met Ala Thr Tyr Asn 

Gly Glu Lys Phe Ile Ser Glu Gln Leu Asp Ser Ile Arg Gln Gln Thr 

Leu Lys Pro Asp Tyr Val Leu Leu Arg Asp Asp Cys Ser Thr Asp Glu 

Thr Val Asn Val Val Asn Asn Tyr Ile Ala Lys His Glu Leu Glu Gly 

Trp Lys Ile Val Lys Asn Asp Lys Asn Leu Gly Trp Arg Leu Asn Phe 

Arg Gln Leu Leu Ile Asp Val Leu Ala Tyr Glu Val Asp Tyr Val Phe 

Phe Ser Asp Gln Asp Asp Ile Trp Tyr Leu Asp Lys Asn Glu Arg Gln 

Phe Ala Ile Met Ser Asp Asn Pro Gln Ile Glu Val Leu Ser Ala Asp 

Val Asp Ile Lys Thr Met Ser Thr Glu Ala Ser Val Pro His Phe Leu 

Thr Phe Ser Ser Asp Arg Ile Ser Gln Tyr Pro Lys Val Tyr Asp 

Tyr Gln Thr Phe Arg Pro Gly Trp Thr Ile Ala Met Lys Arg Asp Phe 



Ala Gln Ala Ile Ala 180

<210> 107

<211> 639

<212> DNA

<213> Streptococcus agalactiae

<400> 107

gtgattatgg ataagtctat tootaaagca actgotaaac gtttatcact gtactaccgt 60 atttttaaac gttttaatac tgatggcate gaaaaagcta gttccaaaca aattgcagat 120 gccctaggta tcgattctgc tactgttcga cgtgatttt cttattttgg tgaactagga 180 cgccgtggtt ttggttatga tgtcaaaaaa cttatgaact tctttgcaga aatattgaac 240 gatcattcta caacaaatgt tatgctggtg gggtgtggaa atatcggtag agctctcttg 300 cattatcgtt tccacgatcg caataaaatg caaatttcaa tggcttttga tttagatagc 360 aatgatttag ttggtaaaac aaccgaggat ggaattcctg tctacggtat ttcgactatc 420 aatgaccatt taatagatag tgatattgaa actgctatcc taacagtacc tagtacagaa 480 gcccaagaag ttgctgacat cttagtcaaa gcaggtataa aaggcatctt gagtttttct 540 ccagttcatt taacattacc aaaagatatc attgttcagt atgtagatt aacaagcgaa 600 ttacaaactt tactttatt catgaaccag cagcgataa

<210> 108

<211> 212

<212> PRT

<213> Streptococcus agalactiae

<400> 108

Met Ile Met Asp Lys Ser Ile Pro Lys Ala Thr Ala Lys Arg Leu Ser

1 5 10 15

Leu Tyr Tyr Arg Ile Phe Lys Arg Phe Asn Thr Asp Gly Ile Glu Lys
20 25 30

Ala Ser Ser Lys Gln Ile Ala Asp Ala Leu Gly Ile Asp Ser Ala Thr

35



117

Val Arg Arg Asp Phe Ser Tyr Phe Gly Glu Leu Gly Arg Arg Gly Phe
50 55 60

Gly Tyr Asp Val Lys Lys Leu Met Asn Phe Phe Ala Glu Ile Leu Asn
65 70 75 80

Asp His Ser Thr Thr Asn Val Met Leu Val Gly Cys Gly Asn Ile Gly
85 90 95

Arg Ala Leu Leu His Tyr Arg Phe His Asp Arg Asn Lys Met Gln Ile 100 105 110

Ser Met Ala Phe Asp Leu Asp Ser Asn Asp Leu Val Gly Lys Thr Thr
115 120 125

Glu Asp Gly Ile Pro Val Tyr Gly Ile Ser Thr Ile Asn Asp His Leu 130 135 140

Ala Gln Glu Val Ala Asp Ile Leu Val Lys Ala Gly Ile Lys Gly Ile
165 170 175

Leu Ser Phe Ser Pro Val His Leu Thr Leu Pro Lys Asp Ile Ile Val 180 185 190

Gln Tyr Val Asp Leu Thr Ser Glu Leu Gln Thr Leu Leu Tyr Phe Met 195 200 205

Asn Gln Gln Arg

210



<210>	109		
<211>	476		
<212>	DNA		

## <213> Streptococcus agalactiae

<400> 109

atgggtgcta aaggagcaga tgtcattctc gttttatcac actctggcat tggagatgat 60 cgatatgaag aaggtgaaga aaacgttggc tatcaaattg ccagcatcaa gggagtggat 120 geegttgtta egggaeaete acaegetgaa ttteeateag gtaaeggtae tggettetat 180 gaaaaataca ctggagttga tggtatcaat ggaaaaataa atggaacacc tgttacaatg 240 gcaggcaagt acggggatca ccttggtatt attgatttag gacttagtta tactaatgga 300 aaatggcaag totocgaaag cagtgctaaa atccgtaaaa ttgatatgaa ctcaacaact 360 gctgacgagc gtatcattgc attggctaag gaagcacacg atggcactat caactatgtt 420 cgccaacaag taggtacaac aactgcgcca attacaagtt actttgcact agttaa

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<210> 110
<211> 158
<212> PRT
<213> Streptococcus agalactiae
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<400> 110

Met Gly Ala Lys Gly Ala Asp Val Ile Leu Val Leu Ser His Ser Gly 15 10 1 5

Ile Gly Asp Asp Arg Tyr Glu Glu Glu Glu Asn Val Gly Tyr Gln 25 30 20

Ile Ala Ser Ile Lys Gly Val Asp Ala Val Val Thr Gly His Ser His 45 40 35

Ala Glu Phe Pro Ser Gly Asn Gly Thr Gly Phe Tyr Glu Lys Tyr Thr 60 55 50

Gly Val Asp Gly Ile Asn Gly Lys Ile Asn Gly Thr Pro Val Thr Met 70 75 80 65



Ala Gly Lys Tyr Gly Asp His Leu Gly Ile Ile Asp Leu Gly Leu Ser 85 90 95

Tyr Thr Asn Gly Lys Trp Gln Val Ser Glu Ser Ser Ala Lys Ile Arg 100 105 110

Lys Ile Asp Met Asn Ser Thr Thr Ala Asp Glu Arg Ile Ile Ala Leu 115 120 125

Ala Lys Glu Ala His Asp Gly Thr Ile Asn Tyr Val Arg Gln Gln Val 130 135 140

Gly Thr Thr Thr Ala Pro Ile Thr Ser Tyr Phe Ala Leu Val 145 150 155

<210> 111

<211> 170

<212> DNA

<213> Streptococcus agalactiae

<400> 111

ttgtcaataa ggtttcaaat cagcttgaaa tatgataaaa taaaacagat tgtaagtgac 60 tgtttaagct tgttttcag agaggtttt atgaatacaa acacaataaa aaaggttgta 120 gcgactggaa ttggagctgc actttttatc attataggta tgctagttaa 170

<210> 112

<211> 56

<212> PRT

<213> Streptococcus agalactiae

<400> 112

Met Ser Ile Arg Phe Gln Ile Ser Leu Lys Tyr Asp Lys Ile Lys Gln
1 5 10 15

Ile Val Ser Asp Cys Leu Ser Leu Phe Phe Arg Glu Val Phe Met Asn

20



120

Thr Asn Thr Ile Lys Lys Val Val Ala Thr Gly Ile Gly Ala Ala Leu
35 40 45

Phe Ile Ile Gly Met Leu Val
50 55

<210> 113

<211> 242

<212> DNA

<213> Streptococcus agalactiae

<400> 113

atgaaacatt taaaatttca atcggtcttc gacattattg gtcctgttat gattggacca 60 tcaagtagtc atactgcagg agctgtccgc attggtaaag ttgtccattc tatttttggt 120 gaacctagtg aagtaacctt tcatttatac aattcttttg ctaaaactta ccaaggacac 180 ggtactgata aagcattggt tgcagggatt ctaggaatgg atacagataa tccagatatt 240 aa 242

<210> 114

<211> 80

<212> PRT

<213> Streptococcus agalactiae

<400> 114

Met Lys His Leu Lys Phe Gln Ser Val Phe Asp Ile Ile Gly Pro Val

1 5 10 15

Met Ile Gly Pro Ser Ser Ser His Thr Ala Gly Ala Val Arg Ile Gly
20 25 30

Lys Val Val His Ser Ile Phe Gly Glu Pro Ser Glu Val Thr Phe His
35 40 45

Leu Tyr Asn Ser Phe Ala Lys Thr Tyr Gln Gly His Gly Thr Asp Lys
50 55 60





Ala Leu Val Ala Gly Ile Leu Gly Met Asp Thr Asp Asn Pro Asp Ile 65 70 75 80

<2:	۷٥>	11	5
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<211> 122

<212> DNA

<213> Streptococcus agalactiae

## <400> 115

gtgtcagaag gtgttttaat gtttctaaaa gaagatgacg tagagacttt tcttcatatc 60 ctgacaaatt catttagcca atttatggca caatttgatt tgtgtcataa ggaaatgatt 120 122

<210> 116

<211> 83

<212> DNA

<213> Streptococcus agalactiae

## <400> 116

atgacctaca aagattacac aggtttagat cggactgaac ttttgagtaa agtgcgtcat 60 atgatgtccg acaaacgttt taa 83

<210> 117

<211> 27

<212> PRT

<213> Streptococcus agalactiae

## <400> 117

Met Thr Tyr Lys Asp Tyr Thr Gly Leu Asp Arg Thr Glu Leu Leu Ser 1 5 15



122

Lys Val Arg His Met Met Ser Asp Lys Arg Phe
20 25

<210> 118

<211> 94

<212> DNA

<213> Streptococcus agalactiae

<400> 118

ctgagttggg tcttggaaac ggtcctgtca atcatactag ctatcaagga gactaaaatg 60 tatttagaac aactaaaaga ggtaaatcct ttaa 94

<210> 119

<211> 31

<212> PRT

<213> Streptococcus agalactiae

<400> 119

Met Ser Trp Val Leu Glu Thr Val Leu Ser Ile Ile Leu Ala Ile Lys

1 5 10 15

Glu Thr Lys Met Tyr Leu Glu Gln Leu Lys Glu Val Asn Pro Leu
20 25 30

<210> 120

<211> 1230

<212> DNA

<213> Streptococcus agalactiae

<400> 120

gtgaaaaaa aattagtoto atoacttota aagtgttoto taatoattat tgttagottt 60 gctggtggag catttgctag ttttgtcatg aatcataatg acaatattoo aaatggtggt 120 gtcactaaaa ctagtaaagt aaattataat aacataacgo ctacaacaaa agctgttaaa 180 aaggtacaaa atagtgttgt ttctgttato aattataac aacaagagag tcgttctgac 240







ctatcagact tctatagtca tttttttggt aatcaggggg gcaacactga taagggctta 300 caagtttacg gtgaaggctc tggagtcatc tataaaaaag atggtaaaaa tgcctatgtt 360 gtcactaata accacgtcat tgatggggct aaacaaattg aaattcaact agctgatggc 420 tcaaaagcag ttgggaaact tgttgggtca gatacctact ctgatttagc cgtcgtcaaa 480 attocatoag ataaagttto aaatattgoa gaatttgotg attoatoaaa actoaacatt 540 ggtgaaactg ctatagcgat cggaagccct cttggaactg agtatgcaaa ttctgtaact 600 caaggtattg tatctagttt aaaaagaact gtaacaatga ctaatgaaga aggacaaaca 660 qtttctacaa atgctatcca gacggatgct gctatcaatc ctggtaattc aggtggagca 720 cttatcaata ttgaaggaca ggttattgga attaattcta gtaaaatttc ttctacatca 780 aatcaaacct caggacaatc gtcaggaaat agcgttgaag gtatgggatt tgccattcct 840 tcaaatgatg ttgttaagat tatcaatcaa cttgagagta acggacaagt agagagacct 900 gctctaggta tttctatggc tggattaagt aatttaccat ccgatgttat tagtaaactg 960 aaaatcccaa gtaatgttac taatggtatt gtagtagcat ctatccaatc tggcatgcca 1020 gctcaaggca aactaaagaa atacgatgtc attactaaag ttgacgataa agaagtagca 1080 tctccaaqtg atttacaaag tttactctat ggccaccagg taggggattc cataacagta 1140 accttttatc gtggtgaaaa taaacaaaca gtcactataa aacttactaa aactagtaaa 1200

123

<210> 121

<211> 409

<212> PRT

<213> Streptococcus agalactiae

gatttagcta aacaacgagc aaataactaa

<400> 121

Met Lys Lys Lys Leu Val Ser Ser Leu Leu Lys Cys Ser Leu Ile Ile 15 5 10 1

Ile Val Ser Phe Ala Gly Gly Ala Phe Ala Ser Phe Val Met Asn His 30 20 25

Asn Asp Asn Ile Pro Asn Gly Gly Val Thr Lys Thr Ser Lys Val Asn 35 40 45

Tyr Asn Asn Ile Thr Pro Thr Thr Lys Ala Val Lys Lys Val Gln Asn 50 55 60

124

Ser Val Val Ser Val Ile Asn Tyr Lys Gln Glu Ser Arg Ser Asp
65 70 75 80

Leu Ser Asp Phe Tyr Ser His Phe Phe Gly Asn Gln Gly Gly Asn Thr
85 90 95

Asp Lys Gly Leu Gln Val Tyr Gly Glu Gly Ser Gly Val Ile Tyr Lys 100 105 110

Lys Asp Gly Lys Asn Ala Tyr Val Val Thr Asn Asn His Val Ile Asp 115 120 125

Gly Ala Lys Gln Ile Glu Ile Gln Leu Ala Asp Gly Ser Lys Ala Val 130 135 140

Gly Lys Leu Val Gly Ser Asp Thr Tyr Ser Asp Leu Ala Val Val Lys
145 150 155 160

Ile Pro Ser Asp Lys Val Ser Asn Ile Ala Glu Phe Ala Asp Ser Ser 165 170 175

Lys Leu Asn Ile Gly Glu Thr Ala Ile Ala Ile Gly Ser Pro Leu Gly
180 185 190

Thr Glu Tyr Ala Asn Ser Val Thr Gln Gly Ile Val Ser Ser Leu Lys
195 200 205

Arg Thr Val Thr Met Thr Asn Glu Glu Gly Gln Thr Val Ser Thr Asn 210 215 220

Ala Ile Gln Thr Asp Ala Ala Ile Asn Pro Gly Asn Ser Gly Gly Ala 225 230 235 240

Leu Ile Asn Ile Glu Gly Gln Val Ile Gly Ile Asn Ser Ser Lys Ile
245 250 255



125

Ser Ser Thr Ser Asn Gln Thr Ser Gly Gln Ser Ser Gly Asn Ser Val
260 265 270

Glu Gly Met Gly Phe Ala Ile Pro Ser Asn Asp Val Val Lys Ile Ile
275 280 285

Asn Gln Leu Glu Ser Asn Gly Gln Val Glu Arg Pro Ala Leu Gly Ile 290 295 300

Ser Met Ala Gly Leu Ser Asn Leu Pro Ser Asp Val Ile Ser Lys Leu 305 310 315 320

Lys Ile Pro Ser Asn Val Thr Asn Gly Ile Val Val Ala Ser Ile Gln 325 330 335

Ser Gly Met Pro Ala Gln Gly Lys Leu Lys Lys Tyr Asp Val Ile Thr 340 345 350

Lys Val Asp Asp Lys Glu Val Ala Ser Pro Ser Asp Leu Gln Ser Leu 355 360 365

Leu Tyr Gly His Gln Val Gly Asp Ser Ile Thr Val Thr Phe Tyr Arg 370 375 380

Gly Glu Asn Lys Gln Thr Val Thr Ile Lys Leu Thr Lys Thr Ser Lys 385 390 395 400

Asp Leu Ala Lys Gln Arg Ala Asn Asn
405

<210> 122

<211> 1923

<212> DNA

<213> Streptococcus agalactiae

<400> 122

atgttaaaat ggtatacaaa caaaggaggg aggatgataa tgaagaaatg ttttttggct 60 atttgtttag ctcttagttt ttttatggtt tcagttcaag cagatgaggt ggactataac 120







attcctcatt	atgagggtaa	tctaactatt	cacaatgata	atagtgctga	ttttacagag	180
aaggttactt	accaatttga	ttcgtcctat	aatggacagt	atgtcacgtt	aggtacggcg	240
ggtaagttat	ctgacaattt	tgatattaat	aataagccac	aggttgaagt	ttcaattaat	300
ggtaaagtaa	ggaaagttag	ttaccagata	gaagatttgg	aggatggcta	ccgtttgaaa	360
gtgtttaatg	gtggtgaagc	aggtgatact	gttaaagtca	atgttcagtg	gaaactaaaa	420
aatgttctat	ttatgcataa	ggatgttggt	gaacttaact	ggattcctat	tagcgactgg	480
gataaaacgt	tagagaaagt	agatttttgg	atatcaactg	acaaaaaggt	tgctctttct	540
cgtctttggg	ggcacttggg	ttatcttaaa	actcctccta	aaataagaca	aaataataat	600
cgttaccatt	tgacagcttt	taatgtaaac	aaacgattag	aatttcatgg	ttattgggat	660
agatcttatt	ttaatctacc	tacaaacagt	aaaaataatt	acaagaaaaa	aattgaacat	720
caagagaaga	taatagagcg	tcatggtttt	atcctaagtt	tcttgttaag	gatattatta	780
ccttcattct	ttattattgt	gacactattc	atctcaatta	gggtgttcct	gtttagaaaa	840
aaagttaata	aatacgggca	attccctaag	gatcatcatt	tatatgaagc	acctgaggac	900
ctttcaccat	tagagttaac	tcaaagcatt	tatagtatga	gctttaaaaa	ttttcaagat	960
gaggagaaga	aaactcacct	tatcagtcaa	gaacaactca	tacagtcaat	tctattagac	1020
ttgattgata	gaaaagtatt	gaattatgat	gataacttgt	tatctctagc	taacttagat	1080
agagcttctg	atgcagaaat	agattttata	gagtttgctt	ttgcggattc	tacgagtttg	1140
aagccagatc	aactcttttc	taattaccaa	tttagttata	aagaaacact	acgtgaactg	1200
aaaaagcagc	acaaggette	agatctgcaa	aatcaaatga	gacgccgagg	aagtaatgcc	1260
ttatcaagaa	ttacgcgtct	cacaaggttg	atttctaaag	acaatataaa	ctctcttaga	1320
agaaagggaa	tttcatcccc	ttatcgtaaa	atgtcttcag	aagagtctaa	agaattatct	1380
aggttaaaaa	gattcagtta	cctatcacct	cttatttctt	ttgttgttat	aatttatacg	1440
cttttttaa	attattttac	ctatttctgt	atctatctct	tattgtttgg	tgttatcctg	1500
ttgttgaata	aaatcatttt	tatgatgaca	agaaaaataa	gtaacggtta	tattgtaact	1560
gaagatggag	caagtcgtgt	ctaccaatgg	actagtttta	ggaacatgct	aagggatatc	1620
aaatcgtttg	atcgttcaga	gttagaaagt	atcgtattat	ggaatcgaat	attggtttac	1680
gctactttat	teggetaege	tgaccgtgtt	gagaaagtac	tcagagtgaa	ccaaatagat	1740
attccagaaa	gatttgcaaa	cattgatagt	catcgatttg	cgatttcagt	caatcaatct	1800
					tagtgttaat	
tcaggcggtt	cttcaggtgg	tttctcaggc	ggcggaggcg	gcggaggtgg	cggtgccttc	1920
taa						1923

<210> 123

<211> 640

<212> PRT

<213> Streptococcus agalactiae



127

<400> 123

Met Leu Lys Trp Tyr Thr Asn Lys Gly Gly Arg Met Ile Met Lys Lys

1 5 10 15

Cys Phe Leu Ala Ile Cys Leu Ala Leu Ser Phe Phe Met Val Ser Val
20 25 30

Gln Ala Asp Glu Val Asp Tyr Asn Ile Pro His Tyr Glu Gly Asn Leu
35 40 45

Thr Ile His Asn Asp Asn Ser Ala Asp Phe Thr Glu Lys Val Thr Tyr
50 55 60

Gln Phe Asp Ser Ser Tyr Asn Gly Gln Tyr Val Thr Leu Gly Thr Ala
65 70 75 80

Gly Lys Leu Ser Asp Asn Phe Asp Ile Asn Asn Lys Pro Gln Val Glu 85 90 95

Val Ser Ile Asn Gly Lys Val Arg Lys Val Ser Tyr Gln Ile Glu Asp 100 105 110

Leu Glu Asp Gly Tyr Arg Leu Lys Val Phe Asn Gly Gly Glu Ala Gly
115 120 125

Asp Thr Val Lys Val Asn Val Gln Trp Lys Leu Lys Asn Val Leu Phe 130 135 140

Met His Lys Asp Val Gly Glu Leu Asn Trp Ile Pro Ile Ser Asp Trp 145 150 155 160

Asp Lys Thr Leu Glu Lys Val Asp Phe Trp Ile Ser Thr Asp Lys Lys
165 170 175

Val Ala Leu Ser Arg Leu Trp Gly His Leu Gly Tyr Leu Lys Thr Pro 180 185 190



128

Pro Lys Ile Arg Gln Asn Asn Asn Arg Tyr His Leu Thr Ala Phe Asn 195 200 205

Val Asn Lys Arg Leu Glu Phe His Gly Tyr Trp Asp Arg Ser Tyr Phe 210 215 220

Asn Leu Pro Thr Asn Ser Lys Asn Asn Tyr Lys Lys Lys Ile Glu His 225 230 235 240

Gln Glu Lys Ile Ile Glu Arg His Gly Phe Ile Leu Ser Phe Leu Leu 245 250 255

Arg Ile Leu Leu Pro Ser Phe Phe Ile Ile Val Thr Leu Phe Ile Ser 260 265 270

Ile Arg Val Phe Leu Phe Arg Lys Lys Val Asn Lys Tyr Gly Gln Phe
275 280 285

Pro Lys Asp His His Leu Tyr Glu Ala Pro Glu Asp Leu Ser Pro Leu 290 295 300

Glu Leu Thr Gln Ser Ile Tyr Ser Met Ser Phe Lys Asn Phe Gln Asp 305 310 315 320

Glu Glu Lys Lys Thr His Leu Ile Ser Gln Glu Gln Leu Ile Gln Ser 325 330 335

Ile Leu Leu Asp Leu Ile Asp Arg Lys Val Leu Asn Tyr Asp Asp Asn 340 345 350

Leu Leu Ser Leu Ala Asn Leu Asp Arg Ala Ser Asp Ala Glu Ile Asp 355 360 365

Phe Ile Glu Phe Ala Phe Ala Asp Ser Thr Ser Leu Lys Pro Asp Gln 370 375 380

Leu Phe Ser Asn Tyr Gln Phe Ser Tyr Lys Glu Thr Leu Arg Glu Leu 385 390 395 400



129

Lys Lys Gln His Lys Ala Ser Asp Leu Gln Asn Gln Met Arg Arg Arg
405 410 415

Gly Ser Asn Ala Leu Ser Arg Ile Thr Arg Leu Thr Arg Leu Ile Ser
420 425 430

Lys Asp Asn Ile Asn Ser Leu Arg Arg Lys Gly Ile Ser Ser Pro Tyr
435 440 445

Arg Lys Met Ser Ser Glu Glu Ser Lys Glu Leu Ser Arg Leu Lys Arg 450 455 460

Phe Ser Tyr Leu Ser Pro Leu Ile Ser Phe Val Val Ile Ile Tyr Thr 465 470 475 480

Leu Phe Leu Asn Tyr Phe Thr Tyr Phe Cys Ile Tyr Leu Leu Phe
485 490 495

Gly Val Ile Leu Leu Leu Asn Lys Ile Ile Phe Met Met Thr Arg Lys
500 505 510

Ile Ser Asn Gly Tyr Ile Val Thr Glu Asp Gly Ala Ser Arg Val Tyr
515 520 525

Gln Trp Thr Ser Phe Arg Asn Met Leu Arg Asp Ile Lys Ser Phe Asp 530 535 540

Arg Ser Glu Leu Glu Ser Ile Val Leu Trp Asn Arg Ile Leu Val Tyr 545 550 555 560

Ala Thr Leu Phe Gly Tyr Ala Asp Arg Val Glu Lys Val Leu Arg Val
565 570 575

Asn Gln Ile Asp Ile Pro Glu Arg Phe Ala Asn Ile Asp Ser His Arg
580 585 590

Phe Ala Ile Ser Val Asn Gln Ser Ser Asn His Phe Ser Thr Ile Thr
595 600 605





Glu Asp Val Ser His Ala Ser Asn Phe Ser Val Asn Ser Gly Gly Ser 610 615 620

Ser Gly Gly Phe Ser Gly Gly Gly Gly Gly Gly Gly Gly Ala Phe 625 630 635 640

<210> 124

<211> 2712

<212> DNA

<213> Streptococcus agalactiae

<400> 124

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aattottttt taaataaaat accagataga agtggtatto tocaagagga ttttataato 1320 aatggtgatg attatcaaat agtaaaagga gatggagaga gttttaaact gttttcggat 1380 agaaaagtto otgttactgg aggaacgaca caagcagott atcgagtaco gcaaaatcaa 1440 ctctctgtaa tgagtaatga gggatatgca attaatagtg gatatattta tctctattgg 1500 agagattaca actgggtcta tccatttgat cctaagacaa agaaagtttc tgcaacgaaa 1560 caaatcaaaa ctcatggtga gccaacaaca ttatacttta atggaaatat aagacctaaa 1620 qqttatqaca tttttactgt tgggattggt gtaaacggag atcctggtgc aactcctctt 1680 gaagctgaga aatttatgca atcaatatca agtaaaacag aaaattatac taatgttgat 1740 qatacaaata aaatttatga tgagctaaat aaatacttta aaacaattgt tgaggaaaaa 1800 cattctattg ttgatggaaa tgtgactgat cctatgggag agatgattga attccaatta 1860 aaaaatqqtc aaaqttttac acatgatgat tacgttttgg ttggaaatga tggcagtcaa 1920 ttaaaaaatg gtgtggctct tggtggacca aacagtgatg ggggaatttt aaaagatgtt 1980 acagtgactt atgataagac atctcaaacc atcaaaatca atcatttgaa cttaggaagt 2040 qqacaaaaag tagttcttac ctatgatgta cgtttaaaag ataactatat aagtaacaaa 2100 ttttacaata caaataatcg tacaacgcta agtccgaaga gtgaaaaaga accaaatact 2160 attogtgatt toccaattoo caaaattogt gatgttogtg agtttooggt actaaccato 2220 agtaatcaga agaaaatggg tgaggttgaa tttattaaag ttaataaaga caaacattca 2280 qaatcgcttt tgggagctaa gtttcaactt cagatagaaa aagatttttc tgggtataag 2340 caatttgttc cagagggaag tgatgttaca acaaagaatg atggtaaaat ttatttaaa 2400 gcacttcaag atggtaacta taaattatat gaaatttcaa gtccagatgg ctatatagag 2460 gttaaaacga aacctgttgt gacatttaca attcaaaatg gagaagttac gaacctgaaa 2520 gcagatccaa atgctaataa aaatcaaatc gggtatcttg aaggaaatgg taaacatctt 2580 attaccaaca ctcccaaacg cccaccaggt gtttttccta aaacaggggg aattggtaca 2640 attgtctata tattagttgg ttctactttt atgatactta ccatttgttc tttccgtcgt 2700

131

<210> 125

aaacaattgt aa

<211> 903

<212> PRT

<213> Streptococcus agalactiae

20

<400> 125

Met Met Ile Val Asn Asn Gly Tyr Leu Glu Gly Arg Lys Met Lys Lys

1 5 10 15

Arg Gln Lys Ile Trp Arg Gly Leu Ser Val Thr Leu Leu Ile Leu Ser

25

30

2712

Gln Ile Pro Phe Gly Ile Leu Val Gln Gly Glu Thr Gln Asp Thr Asn Gln Ala Leu Gly Lys Val Ile Val Lys Lys Thr Gly Asp Asn Ala Thr Pro Leu Gly Lys Ala Thr Phe Val Leu Lys Asn Asp Asn Asp Lys Ser Glu Thr Ser His Glu Thr Val Glu Gly Ser Gly Glu Ala Thr Phe Glu Asn Ile Lys Pro Gly Asp Tyr Thr Leu Arg Glu Glu Thr Ala Pro Ile Gly Tyr Lys Lys Thr Asp Lys Thr Trp Lys Val Lys Val Ala Asp Asn Gly Ala Thr Ile Ile Glu Gly Met Asp Ala Asp Lys Ala Glu Lys Arg Lys Glu Val Leu Asn Ala Gln Tyr Pro Lys Ser Ala Ile Tyr Glu Asp Thr Lys Glu Asn Tyr Pro Leu Val Asn Val Glu Gly Ser Lys Val Gly Glu Gln Tyr Lys Ala Leu Asn Pro Ile Asn Gly Lys Asp Gly Arg Arg Glu Ile Ala Glu Gly Trp Leu Ser Lys Lys Asn Pro Gly Val Asn Asp Leu Asp Lys Asn Lys Tyr Lys Ile Glu Leu Thr Val Glu Gly Lys Thr 

Thr Val Glu Thr Lys Glu Leu Asn Gln Pro Leu Asp Val Val Leu





 Leu
 Asp
 Asn
 Ser
 Asn
 Ser
 Met
 Asn
 Asn
 Glu
 Arg
 Ala
 Asn
 Asn
 Ser
 Gln

 Arg
 Ala
 Leu
 Lys
 Ala
 Glu
 Ala
 Glu
 Lys
 Leu
 Lys
 Lys
 Lys
 Ile

 Arg
 Ala
 Leu
 Lys
 Ala
 Glu
 Ala
 Val
 Glu
 Lys
 Leu
 Lys
 Lys
 Lys
 Ile

 Arg
 Lys
 L

Thr Ser Asn Lys Asp Asn Arg Val Ala Leu Val Thr Tyr Ala Ser Thr
275 280 285

Ile Phe Asp Gly Thr Glu Ala Thr Val Ser Lys Gly Val Ala Asp Gln 290 295 300

Asn Gly Lys Ala Leu Asn Asp Ser Val Ser Trp Asp Tyr His Lys Thr 305 310 315 320

Thr Phe Thr Ala Thr Thr His Asn Tyr Ser Tyr Leu Asn Leu Thr Asn
325 330 335

Asp Ala Asn Glu Val Asn Ile Leu Lys Ser Arg Ile Pro Lys Glu Ala 340 345 350

Glu His Ile Asn Gly Asp Arg Thr Leu Tyr Gln Phe Gly Ala Thr Phe 355 360 365

Thr Gln Lys Ala Leu Met Lys Ala Asn Glu Ile Leu Glu Thr Gln Ser 370 375 380

Ser Asn Ala Arg Lys Leu Ile Phe His Val Thr Asp Gly Val Pro 385 390 395 400

Thr Met Ser Tyr Ala Ile Asn Phe Asn Pro Tyr Ile Ser Thr Ser Tyr 405 410 415

Gln Asn Gln Phe Asn Ser Phe Leu Asn Lys Ile Pro Asp Arg Ser Gly
420 425 430

Ile Leu Gln Glu Asp Phe Ile Ile Asn Gly Asp Asp Tyr Gln Ile Val
435 440 445





Lys Gly Asp Gly Glu Ser Phe Lys Leu Phe Ser Asp Arg Lys Val Pro Val Thr Gly Gly Thr Thr Gln Ala Ala Tyr Arg Val Pro Gln Asn Gln Leu Ser Val Met Ser Asn Glu Gly Tyr Ala Ile Asn Ser Gly Tyr Ile Tyr Leu Tyr Trp Arg Asp Tyr Asn Trp Val Tyr Pro Phe Asp Pro Lys Thr Lys Lys Val Ser Ala Thr Lys Gln Ile Lys Thr His Gly Glu Pro Thr Thr Leu Tyr Phe Asn Gly Asn Ile Arg Pro Lys Gly Tyr Asp Ile Phe Thr Val Gly Ile Gly Val Asn Gly Asp Pro Gly Ala Thr Pro Leu Glu Ala Glu Lys Phe Met Gln Ser Ile Ser Ser Lys Thr Glu Asn Tyr Thr Asn Val Asp Asp Thr Asn Lys Ile Tyr Asp Glu Leu Asn Lys Tyr Phe Lys Thr Ile Val Glu Glu Lys His Ser Ile Val Asp Gly Asn Val Thr Asp Pro Met Gly Glu Met Ile Glu Phe Gln Leu Lys Asn Gly Gln Ser Phe Thr His Asp Asp Tyr Val Leu Val Gly Asn Asp Gly Ser Gln 

Leu Lys Asn Gly Val Ala Leu Gly Gly Pro Asn Ser Asp Gly Gly Ile 



Leu Lys Asp Val Thr Val Thr Tyr Asp Lys Thr Ser Gln Thr Ile Lys
660 665 670

Ile Asn His Leu Asn Leu Gly Ser Gly Gln Lys Val Val Leu Thr Tyr
675 680 685

Asp Val Arg Leu Lys Asp Asn Tyr Ile Ser Asn Lys Phe Tyr Asn Thr
690 695 700

Asn Asn Arg Thr Thr Leu Ser Pro Lys Ser Glu Lys Glu Pro Asn Thr
705 710 715 720

Ile Arg Asp Phe Pro Ile Pro Lys Ile Arg Asp Val Arg Glu Phe Pro
725 730 735

Val Leu Thr Ile Ser Asn Gln Lys Lys Met Gly Glu Val Glu Phe Ile 740 745 750

Lys Val Asn Lys Asp Lys His Ser Glu Ser Leu Leu Gly Ala Lys Phe
755 760 765

Gln Leu Gln Ile Glu Lys Asp Phe Ser Gly Tyr Lys Gln Phe Val Pro
770 780

Glu Gly Ser Asp Val Thr Thr Lys Asn Asp Gly Lys Ile Tyr Phe Lys
785 790 795 800

Ala Leu Gln Asp Gly Asn Tyr Lys Leu Tyr Glu Ile Ser Ser Pro Asp 805 810 815

Gly Tyr Ile Glu Val Lys Thr Lys Pro Val Val Thr Phe Thr Ile Gln 820 825 830

Asn Gly Glu Val Thr Asn Leu Lys Ala Asp Pro Asn Ala Asn Lys Asn 835 840 845

Gln Ile Gly Tyr Leu Glu Gly Asn Gly Lys His Leu Ile Thr Asn Thr 850 855 860





Pro Lys Arg Pro Pro Gly Val Phe Pro Lys Thr Gly Gly Ile Gly Thr 865 870 875 880

Ile Val Tyr Ile Leu Val Gly Ser Thr Phe Met Ile Leu Thr Ile Cys
885 890 895

Ser Phe Arg Arg Lys Gln Leu 900

<210> 126

<211> 1251

<212> DNA

<213> Streptococcus agalactiae

<400> 126

atgaatagaa aagttgagga aaaaatggct gggaatcgta ataacgatat gaatgtctat 60 tgttcatttt gtggcaaaag ccaagatgaa gtaaaaaaaa ttattgcagg taatggtgtt 120 ttcatttgta atgaatgtgt ggccttatca caagaaatta ttaaggaaga attagctgag 180 gaagtactgg ctcatttagc agaagtacca aaacctaagg aactattaga aatattaaat 240 caatatgttg tagggcaaga tcgtgctaaa cgtgctttag cagttgctgt ctacaatcat 300 tacaagcgtg ttagttatac cgagagtagt gacgatgatg tagatttgca aaaatccaac 360 attttgatga ttggtccaac tggctcagga aaaaccttct tagcacaaac actggctaaa 420 agocttaatg taccgtttgc tattgcagat gcgacttcat tgaccgaagc aggatacgtt 480 ggagaagatg ttgagaatat tcttcttaaa ttgattcaag ctgctgatta taatgtcgaa 540 cgtgctgagc gtggtattat ctacgttgat gaaatagata aaattgctaa gaaaggcgaa 600 aatgtttcta tcacacgtga tgtgtctggt gaaggtgtac agcaagccct tcttaaaatt 660 attgagggta cggtagcaag tgttccccca cagggtgggc gtaaacatcc taaccaagaa 720 atgattcaaa ttaataccaa gaacatcctt tttattgtcg gtggtgcttt tgatggtatt 780 gaagaccttg tgaagcaacg tttaggcgaa aaagttattg gttttggaca gacaagccgt 840 aaaattgatg acaacgcttc ttatatgcaa gagataattt ctgaggatat tcaaaagttt 900 ggactgattc cagagtttat tggccgttta ccagtagttg cagcgttaga acttcttact 960 gcagaagatc tggttcgtat tctgacagaa ccacgcaatg ctttggttaa acaataccaa 1020 accttattat cttatgatgg tgtagaattg gaatttgacc aggatgctct attggctatc 1080 gctgataagg ctatcgagcg caagactggt gcacgtggtt tacgttctat tattgaagaa 1140 acgatgcttg atatcatgtt tgaaattcca agccaagaag atgtaacaaa agttcgtatc 1200 1251 acaaaqqctg ctgttgaggg tactgacaag cctgttttag agacggctta g

<210> 127
<211> 416
<212> PRT
<213> Streptococcus agalactiae
<400> 127

Met Asn Arg Lys Val Glu Glu Lys Met Ala Gly Asn Arg Asn Asn Asp 1 5 10 15

Met Asn Val Tyr Cys Ser Phe Cys Gly Lys Ser Gln Asp Glu Val Lys
20 25 30

Lys Ile Ile Ala Gly Asn Gly Val Phe Ile Cys Asn Glu Cys Val Ala 35 40 45

Leu Ser Gln Glu Ile Ile Lys Glu Glu Leu Ala Glu Glu Val Leu Ala 50 55 60

His Leu Ala Glu Val Pro Lys Pro Lys Glu Leu Leu Glu Ile Leu Asn 65 70 75 80

Gln Tyr Val Val Gly Gln Asp Arg Ala Lys Arg Ala Leu Ala Val Ala 85 90 95

Val Tyr Asn His Tyr Lys Arg Val Ser Tyr Thr Glu Ser Ser Asp Asp
100 105 110

Asp Val Asp Leu Gln Lys Ser Asn Ile Leu Met Ile Gly Pro Thr Gly
115 120 125

Ser Gly Lys Thr Phe Leu Ala Gln Thr Leu Ala Lys Ser Leu Asn Val 130 135 140

Pro Phe Ala Ile Ala Asp Ala Thr Ser Leu Thr Glu Ala Gly Tyr Val 145 150 155 160



138

Gly Glu Asp Val Glu Asn Ile Leu Leu Lys Leu Ile Gln Ala Ala Asp 165 170 175

Tyr Asn Val Glu Arg Ala Glu Arg Gly Ile Ile Tyr Val Asp Glu Ile 180 185 190

Asp Lys Ile Ala Lys Lys Gly Glu Asn Val Ser Ile Thr Arg Asp Val 195 200 205

Ser Gly Glu Gly Val Gln Gln Ala Leu Leu Lys Ile Ile Glu Gly Thr 210 215 220

Val Ala Ser Val Pro Pro Gln Gly Gly Arg Lys His Pro Asn Gln Glu 225 230 235 240

Met Ile Gln Ile Asn Thr Lys Asn Ile Leu Phe Ile Val Gly Gly Ala
245 250 255

Phe Asp Gly Ile Glu Asp Leu Val Lys Gln Arg Leu Gly Glu Lys Val
260 265 270

Ile Gly Phe Gly Gln Thr Ser Arg Lys Ile Asp Asp Asn Ala Ser Tyr
275 280 285

Met Gln Glu Ile Ile Ser Glu Asp Ile Gln Lys Phe Gly Leu Ile Pro 290 295 300

Glu Phe Ile Gly Arg Leu Pro Val Val Ala Ala Leu Glu Leu Leu Thr 305 310 315 320

Ala Glu Asp Leu Val Arg Ile Leu Thr Glu Pro Arg Asn Ala Leu Val 325 330 335

Lys Gln Tyr Gln Thr Leu Leu Ser Tyr Asp Gly Val Glu Leu Glu Phe 340 345 350

Asp Gln Asp Ala Leu Leu Ala Ile Ala Asp Lys Ala Ile Glu Arg Lys 355 360 365





Thr Gly Ala Arg Gly Leu Arg Ser Ile Ile Glu Glu Thr Met Leu Asp 370 375 380

Ile Met Phe Glu Ile Pro Ser Gln Glu Asp Val Thr Lys Val Arg Ile
385 390 395 400

Thr Lys Ala Ala Val Glu Gly Thr Asp Lys Pro Val Leu Glu Thr Ala
405 410 415

<210> 128

<211> 786

<212> DNA

<213> Streptococcus agalactiae

<400> 128

atgaaaagat tacataaact gtttataacc gtaattgcta cattaggtat gttgggggta 60
atgacctttg gtcttccaac gcagccgcaa aacgtaacgc cgatagtaca tgctgatgtc 120
aattcatctg ttgatacgag ccaggaatt caaaataatt taaaaaatgc tattggtaac 180
ctaccattc aatagttaa tggtatttat gaattaaata ataatcagac aaatttaaat 240
gctgatgtca atgttaaagc gtatgttcaa aatacaattg acaatcaaca aagactatca 300
actgctaatg caatgcttga tagaaccatt cgtcaatatc aaaatcgcag agataccact 360
cttcccgatg caaattggaa accattaggt tggcatcaag tagctactaa tgaccattat 420
gggcatgcag tcgacaaggg gcatttaatt gcctatgct tagctggaaa tttcaaaggt 480
tggggatgctt ccgtgtcaaa tcctcaaaat gttgtcacac aaacagctca ttccaaccaa 540
tcaaatcaaa aaatcaatcg tggacaaaat tattatgaaa gcttagttcg taaggcggtt 600
gaccaaaaca aacgtgttcg ttaccgtgta actccattgt accgtaatga tactgattta 660
gttccatttg caatgcacct agaagctaaa tcacaagatg gcacattaga atttaatgtt 720
gctattccaa acacacaagc atcatacact atggattatg caacaggaga aataacacta 780
aattaa



-

140

<210> 129

<211> 261

<212> PRT

<213> Streptococcus agalactiae

<400> 129

Met Lys Arg Leu His Lys Leu Phe Ile Thr Val Ile Ala Thr Leu Gly

1 5 10 15

Met Leu Gly Val Met Thr Phe Gly Leu Pro Thr Gln Pro Gln Asn Val
20 25 30

Thr Pro Ile Val His Ala Asp Val Asn Ser Ser Val Asp Thr Ser Gln
35 40 45

Glu Phe Gln Asn Asn Leu Lys Asn Ala Ile Gly Asn Leu Pro Phe Gln
50 55 60

Tyr Val Asn Gly Ile Tyr Glu Leu Asn Asn Asn Gln Thr Asn Leu Asn 65 70 75 80

Ala Asp Val Asn Val Lys Ala Tyr Val Gln Asn Thr Ile Asp Asn Gln
85 90 95

Gln Arg Leu Ser Thr Ala Asn Ala Met Leu Asp Arg Thr Ile Arg Gln 100 105 110

Tyr Gln Asn Arg Arg Asp Thr Thr Leu Pro Asp Ala Asn Trp Lys Pro

Leu Gly Trp His Gln Val Ala Thr Asn Asp His Tyr Gly His Ala Val 130 135 140

Asp Lys Gly His Leu Ile Ala Tyr Ala Leu Ala Gly Asn Phe Lys Gly 145 150 155 160

Trp Asp Ala Ser Val Ser Asn Pro Gln Asn Val Val Thr Gln Thr Ala 165 170 175



141

His Ser Asn Gln Ser Asn Gln Lys Ile Asn Arg Gly Gln Asn Tyr Tyr 180 185 190

Glu Ser Leu Val Arg Lys Ala Val Asp Gln Asn Lys Arg Val Arg Tyr 195 200 205

Arg Val Thr Pro Leu Tyr Arg Asn Asp Thr Asp Leu Val Pro Phe Ala 210 215 220

Met His Leu Glu Ala Lys Ser Gln Asp Gly Thr Leu Glu Phe Asn Val 225 230 235 240

Ala Ile Pro Asn Thr Gln Ala Ser Tyr Thr Met Asp Tyr Ala Thr Gly
245 250 255

Glu Ile Thr Leu Asn 260

<210> 130

<211> 621

<212> DNA

<213> Streptococcus agalactiae

<400> 130

atgaaaact atcgaaaact tattgtacta ctacttctaa tctttttgc cattttatg 60 ggagcatatg cttacacgca tattgttgaa aaaagatcc taactagcaa tactattgaa 120 aaaactctac ctgtggtaaa tcagattaag cctcaaacca ttaaagaata ccaaaattac 180 ttaactaagg tagctaaacg taatgttctt cctgtagaca ttcctcaggc attaaataat 240 gaaaaggtag aaattactgc tactgatggc atgcaaacat tcacttggaa tgataaaaat 300 aatcctaagc aaaaggttat cttctatgtt catggaggat catatatcca tcaagcttcc 360 gaattacaat atattttgt caataaacta gctaaaaaat tagatgcaaa agttgtcttt 420 cctattacc ctaaagctcc tacatataat tatagtgatg ctatcccaa aattaaaaa 480 ttataccaaa atacattagc tagcgtcaca tctcacaaac agattatcct agtaggtgaa 540 agtgcaggcg gaggccttgc tttaggtatt gctgataacc ttgcacggag catatcaaac 600 aaccaaaaga aattattta a



142

<210> 131

<211> 206

<212> PRT

<213> Streptococcus agalactiae

<400> 131

Met Lys Asn Tyr Arg Lys Leu Ile Val Leu Leu Leu Leu Ile Phe Phe 1 5 10 15

Ala Ile Phe Met Gly Ala Tyr Ala Tyr Thr His Ile Val Glu Lys Arg
20 25 30

Ser Leu Thr Ser Asn Thr Ile Glu Lys Thr Leu Pro Val Val Asn Gln
35 40 45

Ile Lys Pro Gln Thr Ile Lys Glu Tyr Gln Asn Tyr Leu Thr Lys Val
50 55 60

Ala Lys Arg Asn Val Leu Pro Val Asp Ile Pro Gln Ala Leu Asn Asn 65 70 75 80

Glu Lys Val Glu Ile Thr Ala Thr Asp Gly Met Gln Thr Phe Thr Trp

85 90 95

Asn Asp Lys Asn Asn Pro Lys Gln Lys Val Ile Phe Tyr Val His Gly
100 105 110

Gly Ser Tyr Ile His Gln Ala Ser Glu Leu Gln Tyr Ile Phe Val Asn 115 120 125

Lys Leu Ala Lys Lys Leu Asp Ala Lys Val Val Phe Pro Ile Tyr Pro 130 135 140

Lys Ala Pro Thr Tyr Asn Tyr Ser Asp Ala Ile Pro Lys Ile Lys Lys
145 150 155 160

Leu Tyr Gln Asn Thr Leu Ala Ser Val Thr Ser His Lys Gln Ile Ile 165 170 175



143

Leu Val Gly Glu Ser Ala Gly Gly Gly Leu Ala Leu Gly Ile Ala Asp 180 185 190

Asn Leu Ala Arg Ser Ile Ser Asn Asn Gln Lys Lys Leu Phe 195 200 205

<210> 132

<211> 885

<212> DNA

<213> Streptococcus agalactiae

<400> 132

ttgattctaa taacttccta tgggataata tctttatcac aaaaattgag ggaatttatt 60 atgaagttaa aacatattgt cttaggatta gccttaacaa cacttttagg agtcacattt 120 agtaatcaag aagtttcagc aagctcaact tcaagtaaag ttgttaaagt tggtgttatg 180 accttttctg acactgaaaa agcacgttgg gataaaattg aaaagctagt aggtgataaa 240 gctaaaatca aatttacaga atttacagat tatacacaac caaatcaagc gacagccaat 300 aaggatgtgg atattaatgc ctttcaacat tacaatttct tagaaaactg gaataaggaa 360 aataagaaaa acttaattcc acttgaaaag acttacttag ctccaattcg tatctattct 420 gagaaggtaa aatotottaa aaaattgaaa aaaggagoca otattgcaat tocaaatgat 480 gcaacaaatg gtagccgtgc attgtatgtc cttcagtcag caggtttaat caaattgaat 540 gtttctggta agaaggttgc aacagttgct aatatcacat ctaataaaaa ggatattaat 600 attcaggagt tagatgcgag tcaaacacca cgtgcactca aagatgtaga tgcagctatt 660 attaataata catacattga gcaagctaat ttaaaacctt cagatgctat ctttgttgag 720 aaatcagata aaaattcaaa acaatggatt aatatcattg cgggacgtaa aaattggaaa 780 aagcaaaaga acgctaaagc tatccaagct atcttggatg cttatcacac agatgaagtg 840 885 aaaaaagtta tcaaagatac ttcagctgat attccacaat ggtaa

<210> 133

<211> 294

<212> PRT

<213> Streptococcus agalactiae

5

<400> 133

1

Met Ile Leu Ile Thr Ser Tyr Gly Ile Ile Ser Leu Ser Gln Lys Leu



144

Arg Glu Phe Ile Met Lys Leu Lys His Ile Val Leu Gly Leu Ala Leu 20 25 30

Thr Thr Leu Leu Gly Val Thr Phe Ser Asn Gln Glu Val Ser Ala Ser
35 40 45

Ser Thr Ser Ser Lys Val Val Lys Val Gly Val Met Thr Phe Ser Asp
50 55 60

Thr Glu Lys Ala Arg Trp Asp Lys Ile Glu Lys Leu Val Gly Asp Lys
65 70 75 80

Ala Lys Ile Lys Phe Thr Glu Phe Thr Asp Tyr Thr Gln Pro Asn Gln
85 90 95

Ala Thr Ala Asn Lys Asp Val Asp Ile Asn Ala Phe Gln His Tyr Asn
100 105 110

Phe Leu Glu Asn Trp Asn Lys Glu Asn Lys Lys Asn Leu Ile Pro Leu 115 120 125

Glu Lys Thr Tyr Leu Ala Pro Ile Arg Ile Tyr Ser Glu Lys Val Lys 130 135 140

Ser Leu Lys Lys Leu Lys Lys Gly Ala Thr Ile Ala Ile Pro Asn Asp 145 150 155 160

Ala Thr Asn Gly Ser Arg Ala Leu Tyr Val Leu Gln Ser Ala Gly Leu 165 170 175

Ile Lys Leu Asn Val Ser Gly Lys Lys Val Ala Thr Val Ala Asn Ile 180 185 190

Thr Ser Asn Lys Lys Asp Ile Asn Ile Gln Glu Leu Asp Ala Ser Gln
195 200 205

Thr Pro Arg Ala Leu Lys Asp Val Asp Ala Ala Ile Ile Asn Asn Thr 210 215 220



145

Tyr Ile Glu Gln Ala Asn Leu Lys Pro Ser Asp Ala Ile Phe Val Glu 225 230 235 240

Lys Ser Asp Lys Asn Ser Lys Gln Trp Ile Asn Ile Ile Ala Gly Arg 245 250 255

Lys Asn Trp Lys Lys Gln Lys Asn Ala Lys Ala Ile Gln Ala Ile Leu 260 265 270

Asp Ala Tyr His Thr Asp Glu Val Lys Lys Val Ile Lys Asp Thr Ser 275 280 285

Ala Asp Ile Pro Gln Trp 290

<210> 134

<211> 1350

<212> DNA

<213> Streptococcus agalactiae

## <400> 134

atgreaate aatatgatta tategttatt ggtggaggta gtgcaggeag tggtaceget 60 aatagggeag ceatgtatgg ageaaaagte etgttaattg aaggtggaca agtaggtgga 120 acttgtgtta acttaggttg tgtacetaag aaaateatgt ggtatggtge acaagtttet 180 gagacactee ataagtatag tteaggttat ggttttgaag ecaataatet tagttttgat 240 tttactacte taaaagetaa tegegatget taegtgeage ggtetagaea gtegtatgee 300 getaattttg agegtaatgg ggtegaaaag attgatggat ttgetegtt tattgataae 360 catactattg aagtgaatgg teageaatat aaageteete acattactat tggeaacaggt 420 ggacaceete tttaceetga tattattgga agtgaacttg gtgagactte tgatgattt 480 tttggatggg agacettace aaattetaa ttgattgttg gggegggeta tategeggea 540 gaacttgetg gagtggttaa tgaattagge gttgaaacee atettgeat tagaaaagae 600 catacttaet geggatttga tgacatggta acaagtgagg ttatggetga aatggagaa 660 teaggtatet ettacatge taaceatgta ectaaatee ttaaacgega tgaaggtgge 720 aagttgatt ttgaagetga aaatgggaaa acgettgteg ttgategtg tataaggeet 780 ateggeegtg gaccaaatgt agacatgga ettgaaaata eegatattgt tttaaatga 840 aaagattata tcaaaacaga tgaatttgag aatacteetg tagatggeg gtatgetat 900



146

ggagatgtta atgggaaaat tgccttgaca ccggtagcaa ttgcagcag tcgtcgctta 960 tcagaaagac tttttaatca taaagataac gaaaaattag attaccataa tgtaccttca 1020 gttatttta ctcaccctgt aattgggacg gtaggacttt cagaagcagc agctatcgag 1080 caatttggaa aagataatat caaagtctat acatcaactt ttacctctat gtatacggct 1140 gttaccagta atcgccaagc agttaagatg aagctcataa ccctaggaaa agaggaaaaa 1200 gttattggc ttcatggtgt tggttatggt attgatgaa tgattcaagg tttttcagtt 1260 gctatcaaaa tgggggctac taaagcagac tttgatgata ctgttgctat tcacccaact 1320 ggatctgagg aatttgttac aatgcgctaa

<210> 135

<211> 449

<212> PRT

<213> Streptococcus agalactiae

<400> 135

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1 5 10 15

Ser Gly Thr Ala Asn Arg Ala Ala Met Tyr Gly Ala Lys Val Leu Leu 20 25 30

Ile Glu Gly Gly Gln Val Gly Gly Thr Cys Val Asn Leu Gly Cys Val
35 40 45

Pro Lys Lys Ile Met Trp Tyr Gly Ala Gln Val Ser Glu Thr Leu His
50 55 60

Lys Tyr Ser Ser Gly Tyr Gly Phe Glu Ala Asn Asn Leu Ser Phe Asp
65 70 75 80

Phe Thr Thr Leu Lys Ala Asn Arg Asp Ala Tyr Val Gln Arg Ser Arg 85 90 95

Gln Ser Tyr Ala Ala Asn Phe Glu Arg Asn Gly Val Glu Lys Ile Asp 100 105 110

Gly Phe Ala Arg Phe Ile Asp Asn His Thr Ile Glu Val Asn Gly Gln 115 120 125



Gln Tyr Lys Ala Pro His Ile Thr Ile Ala Thr Gly Gly His Pro Leu 130 135 140

Tyr Pro Asp Ile Ile Gly Ser Glu Leu Gly Glu Thr Ser Asp Asp Phe 145 150 155 160

Phe Gly Trp Glu Thr Leu Pro Asn Ser Ile Leu Ile Val Gly Ala Gly
165 170 175

Tyr Ile Ala Ala Glu Leu Ala Gly Val Val Asn Glu Leu Gly Val Glu 180 185 190

Thr His Leu Ala Phe Arg Lys Asp His Ile Leu Arg Gly Phe Asp Asp 195 200 205

Met Val Thr Ser Glu Val Met Ala Glu Met Glu Lys Ser Gly Ile Ser 210 215 220

Leu His Ala Asn His Val Pro Lys Ser Leu Lys Arg Asp Glu Gly Gly
225 230 235 240

Lys Leu Ile Phe Glu Ala Glu Asn Gly Lys Thr Leu Val Val Asp Arg 245 250 255

Val Ile Trp Ala Ile Gly Arg Gly Pro Asn Val Asp Met Gly Leu Glu 260 265 270

Asn Thr Asp Ile Val Leu Asn Asp Lys Asp Tyr Ile Lys Thr Asp Glu 275 280 285

Phe Glu Asn Thr Ser Val Asp Gly Val Tyr Ala Ile Gly Asp Val Asn 290 295 300

Gly Lys Ile Ala Leu Thr Pro Val Ala Ile Ala Ala Gly Arg Arg Leu 305 310 315 320

Ser Glu Arg Leu Phe Asn His Lys Asp Asn Glu Lys Leu Asp Tyr His
325 330 335



148

Asn Val Pro Ser Val Ile Phe Thr His Pro Val Ile Gly Thr Val Gly 340 345 350

Leu Ser Glu Ala Ala Ala Ile Glu Gln Phe Gly Lys Asp Asn Ile Lys 355 360 365

Val Tyr Thr Ser Thr Phe Thr Ser Met Tyr Thr Ala Val Thr Ser Asn 370 375 380

Arg Gln Ala Val Lys Met Lys Leu Ile Thr Leu Gly Lys Glu Glu Lys 385 390 395 400

Val Ile Gly Leu His Gly Val Gly Tyr Gly Ile Asp Glu Met Ile Gln
405 410 415

Gly Phe Ser Val Ala Ile Lys Met Gly Ala Thr Lys Ala Asp Phe Asp
420 425 430

Asp Thr Val Ala Ile His Pro Thr Gly Ser Glu Glu Phe Val Thr Met
435 440 445

Arg

<210> 136

<211> 1317

<212> DNA

<213> Streptococcus agalactiae

<400> 136

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149

gatttgaatg atttatttac agaccaattt attaaggatg tcaataataa gaacatcatt 420 caagetteta agtetggega taaageetae atgtateeaa taagttetge eecattttat 480 atggcgttca ataaaaaaat gcttaaagat gcaggagttt tgaaacttgt aaaagaaggt 540 tggactacta gtgattttga aaaagtacta aaagcactaa aaaataaagg ctatacacca 600 ggttcattct ttgcaaacgg gcaaggagga gatcaaggac cacgtgcatt ttttgctaat 660 ctttataqtq ctccaataac agataaagaa gtaacaaaat ataccactga cactaaaaat 720 tctgtaaaat caatgaaaaa aatagttgaa tggattaaga aaggctactt gatgaatggg 780 tctcagtatg atggctcagc tgacattcaa aacttcgcca atggacaaac tgctttcact 840 atcctatggg ctccagctca accaaaaact caagcaaaat tattagagtc aagtaaagtg 900 gattaccttg aagtgccatt cccatcagaa gatggaaaac cagatttaga ataccttgtt 960 aatggttttg cggtctttaa taataaagat gaaaacaaag taaaagcctc taagaaattt 1020 atcactttta ttgctgatga taaaaaatgg ggaccaaaag atgttatacg tacaggtgct 1080 ttcccaqtta gaacatcatt tggggatctt tataaaggtg ataaacgtat gatgaagatt 1140 tcaaaatgga ctcaatatta ttcaccatat tacaacacta tcgatggatt ttctgaaatg 1200 agaaccttat ggttcccaat ggttcaatct gtatccaatg gtgatgaaaa accagcagat 1260 qctttqaaag actttactca aaaagcaaat gataccatta aaaaagcagc taaataa

<210> 137

<211> 438

<212> PRT

<213> Streptococcus agalactiae

<400> 137

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Glu Lys Asp Lys Val Glu Ile Thr Trp Trp Ala Phe Pro Thr Phe Thr 35 40 45

Gln Glu Lys Ala Lys Asp Gly Val Gly Thr Tyr Glu Lys Lys Val Ile 50 55 60

Lys Ala Phe Glu Lys Lys Asn Pro Asn Ile Lys Val Lys Leu Glu Thr
65 70 75 80



Ile Asp Phe Thr Ser Gly Pro Glu Lys Ile Thr Thr Ala Ile Glu Ala Gly Thr Ala Pro Asp Val Leu Phe Asp Ala Pro Gly Arg Ile Ile Gln Tyr Gly Lys Asn Gly Lys Leu Ala Asp Leu Asn Asp Leu Phe Thr Asp Gln Phe Ile Lys Asp Val Asn Asn Lys Asn Ile Ile Gln Ala Ser Lys Ser Gly Asp Lys Ala Tyr Met Tyr Pro Ile Ser Ser Ala Pro Phe Tyr Met Ala Phe Asn Lys Lys Met Leu Lys Asp Ala Gly Val Leu Lys Leu Val Lys Glu Gly Trp Thr Thr Ser Asp Phe Glu Lys Val Leu Lys Ala Leu Lys Asn Lys Gly Tyr Thr Pro Gly Ser Phe Phe Ala Asn Gly Gln Gly Gly Asp Gln Gly Pro Arg Ala Phe Phe Ala Asn Leu Tyr Ser Ala Pro Ile Thr Asp Lys Glu Val Thr Lys Tyr Thr Thr Asp Thr Lys Asn Ser Val Lys Ser Met Lys Lys Ile Val Glu Trp Ile Lys Lys Gly Tyr

245 250 255

Leu Met Asn Gly Ser Gln Tyr Asp Gly Ser Ala Asp Ile Gln Asn Phe

Ala Asn Gly Gln Thr Ala Phe Thr Ile Leu Trp Ala Pro Ala Gln Pro 275 280 285



Lys Thr Gln Ala Lys Leu Leu Glu Ser Ser Lys Val Asp Tyr Leu Glu 290 . 295 300

Val Pro Phe Pro Ser Glu Asp Gly Lys Pro Asp Leu Glu Tyr Leu Val 305 310 315 320

Asn Gly Phe Ala Val Phe Asn Asn Lys Asp Glu Asn Lys Val Lys Ala 325 330 335

Ser Lys Lys Phe Ile Thr Phe Ile Ala Asp Asp Lys Lys Trp Gly Pro 340 345 350

Lys Asp Val Ile Arg Thr Gly Ala Phe Pro Val Arg Thr Ser Phe Gly 355 360 365

Asp Leu Tyr Lys Gly Asp Lys Arg Met Met Lys Ile Ser Lys Trp Thr 370 375 380

Gln Tyr Tyr Ser Pro Tyr Tyr Asn Thr Ile Asp Gly Phe Ser Glu Met 385 390 395 400

Arg Thr Leu Trp Phe Pro Met Val Gln Ser Val Ser Asn Gly Asp Glu
405 410 415

Lys Pro Ala Asp Ala Leu Lys Asp Phe Thr Gln Lys Ala Asn Asp Thr
420 425 430

Ile Lys Lys Ala Ala Lys
435

<210> 138

<211> 40

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer



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<210> 139
<211> 43
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
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gaagatette eeegggatea caaacagata aeggegtaaa tag
<210> 140
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72237 Description of Michigan Sequence, frimer
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<210> 141
<211> 32
<212> DNA
<213> Artificial Sequence

211>	32
<212>	DNA
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<223>	Description of Artificial Sequence: Primer
	·
<400>	141
egggat	cott atggacotga atcagogttg to



<210> 142	
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catgatatcg gtacctcaag ctcatatcat tgtccggcaa tggtgtgggc tttttttgtt	60
ttagcggata acaatttcac ac	8
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geggateece egggettaat taatgtttaa acaetagteg aagatetege gaatteteet 60

gtgtgaaatt gttatccgct a





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cgcca	gggtt ttcccagtca cgac		
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<210>	146		

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<223> Description of Artificial Sequence: Primer

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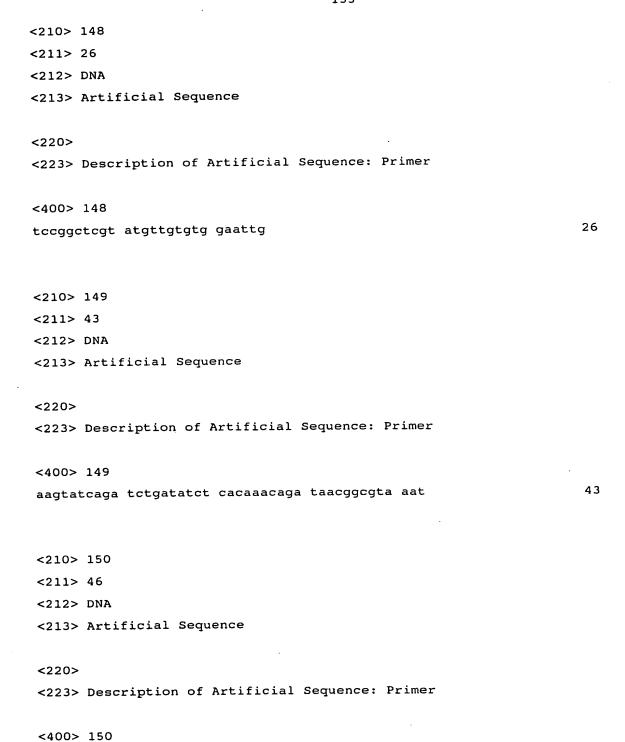
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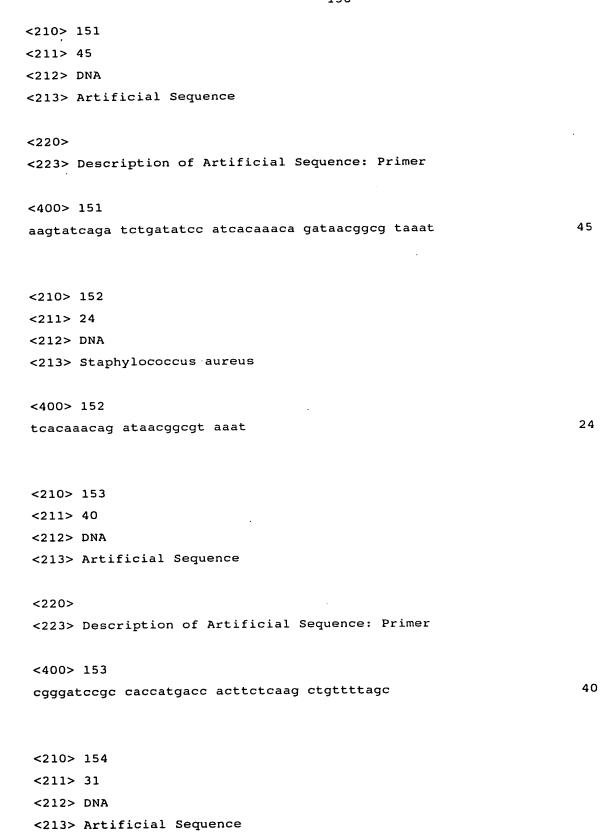
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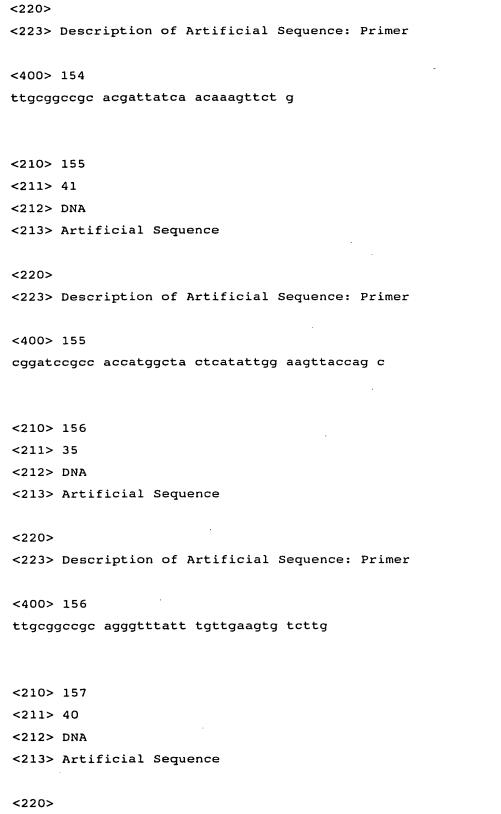




aagtatcaga tottoccogg gatcacaaac agataacggc gtaaat







<223> Description of Artificial Sequence: Primer



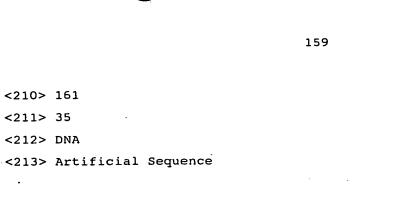




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tigeggeege tecatgeata gaaacageag tece	24
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12207 Description of Artificial Dequence, Filmer	
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<223> Description of Artificial Sequence: Primer

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cggatccgcc accatgattg ttggacacgg aattg 35

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<220>
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<210> 163

<212> DNA

<211> 39

<220>

<213> Artificial Sequence

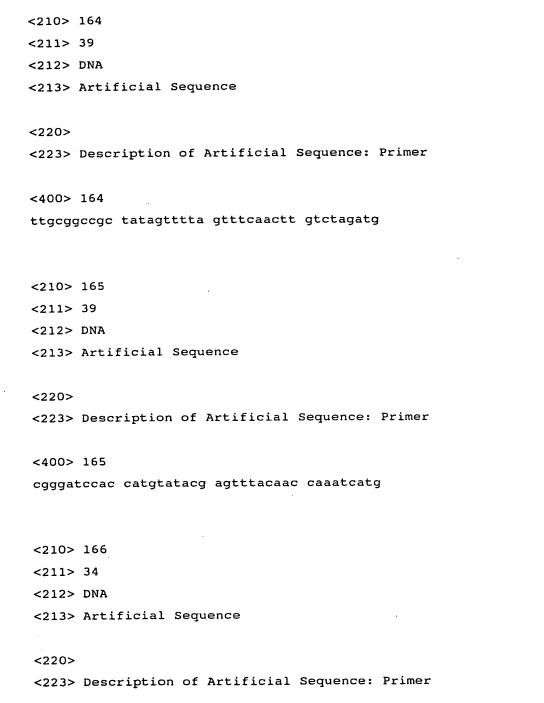
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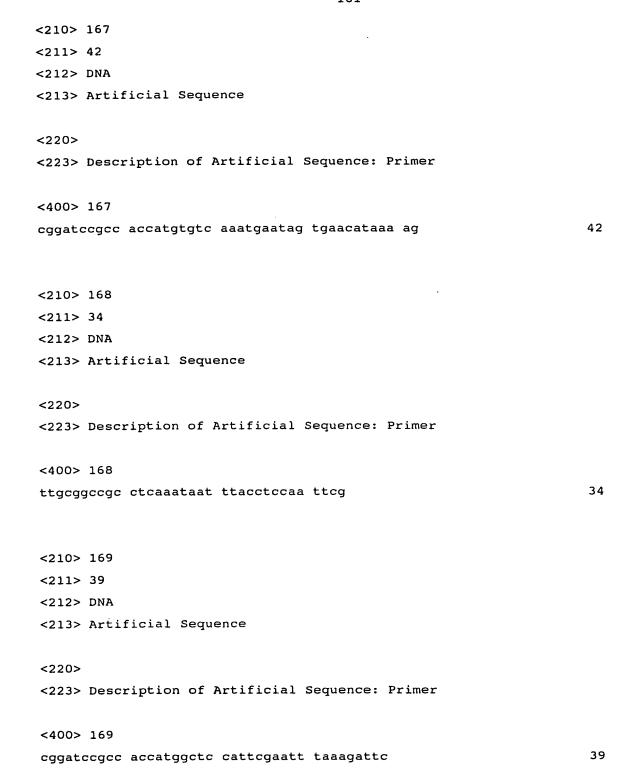
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<400> 166

ttgcggccgc gtcagctcgt actgttttt tagc







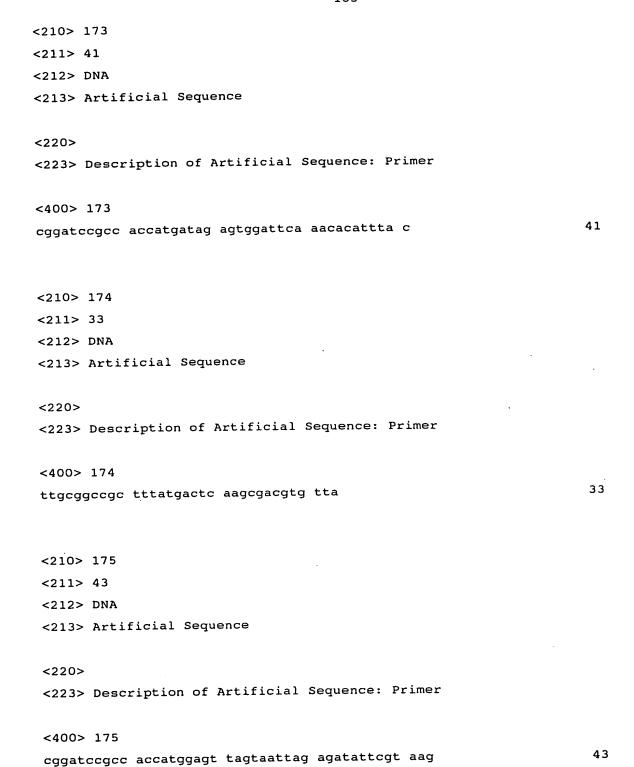
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ttgcggccgc ttctttgttc caactttctg g



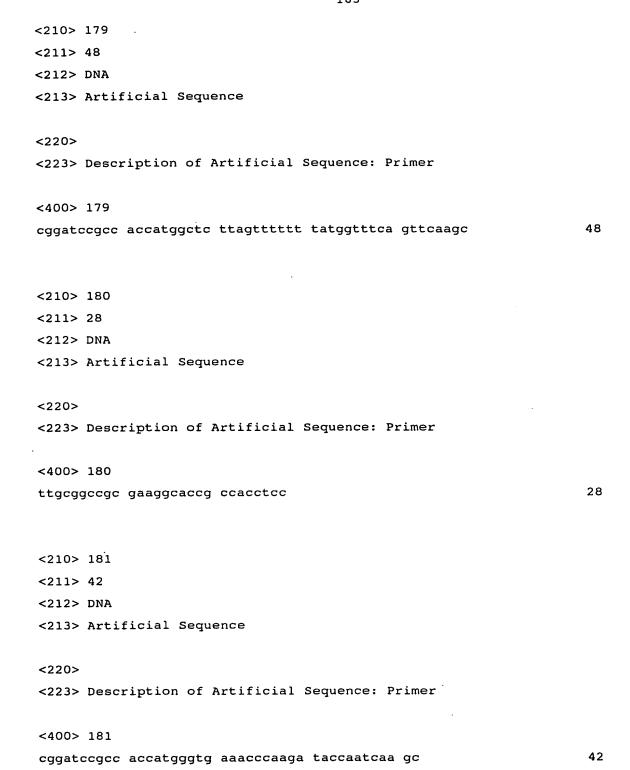


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ttgcggccgc agccgtctct aaaacaggct tg	32

<400> 187

cgggatccac catgtacacg catattgttg aaaaaag





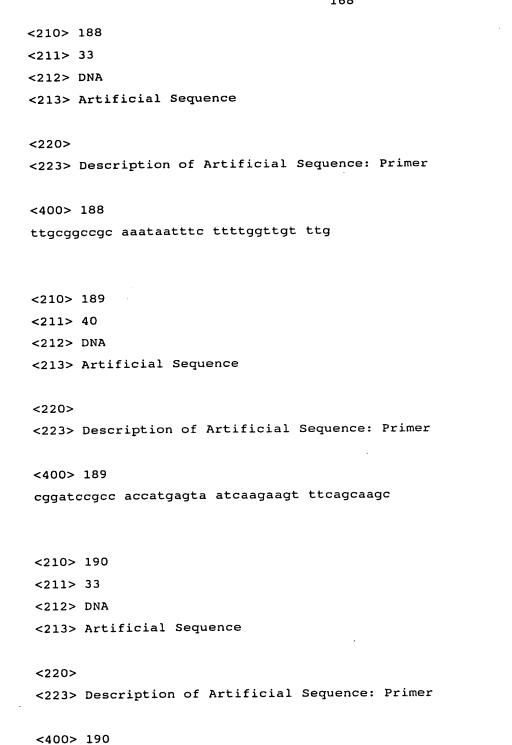
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(213) Altificial bequence
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ttgcggccgc atttagtgtt atttctcctg ttgcataatc c
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37

41







ttgcggccgc ccattgtgga atatcagctg aag



cgggatccac catggctgcc gagaaggata aag



33

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40005		
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<b>\223</b> /	bescription of Artificial bequence. Illinoi	
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	geoge gegeattgta acaaatteet cag 3	3
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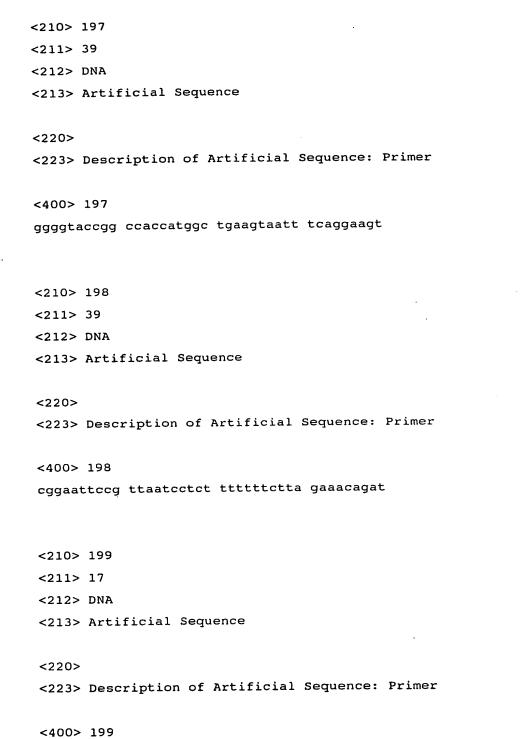




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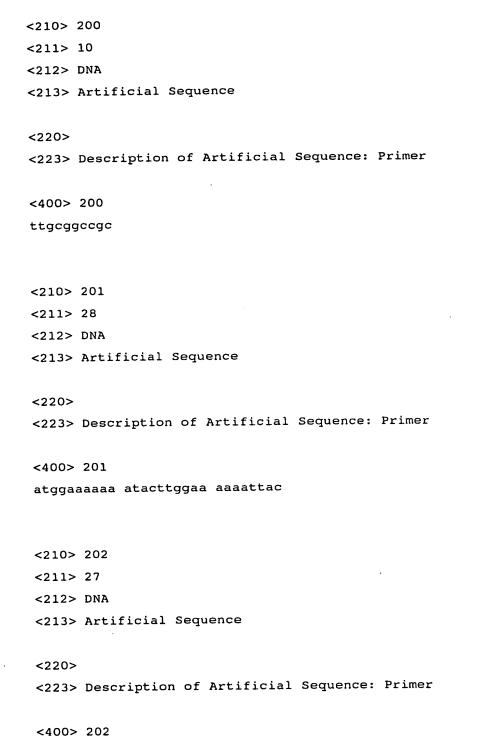
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ctattttgtt ttagcgatgt ctttatc



atgaaaaag ttttttttt catggctatg







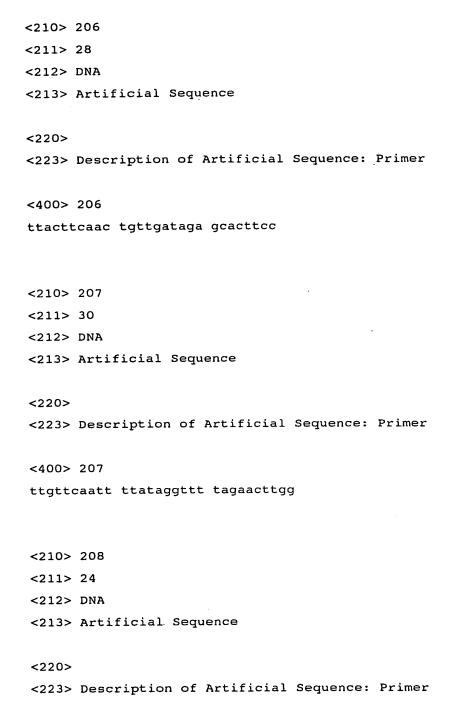
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atgtc	caaaac aaaaagtaac ggcaac	26
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	> 204 ttatgg ccaataccat aagttaattg	30
ttatt	· · · · · · · · · · · · · · · · · · ·	30
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	> DNA	
<213>	> Artificial Sequence	
<220>	)>	
<223>	> Description of Artificial Sequence: Primer	
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<400> 208

ttaattttca ttgcgtctca aacc





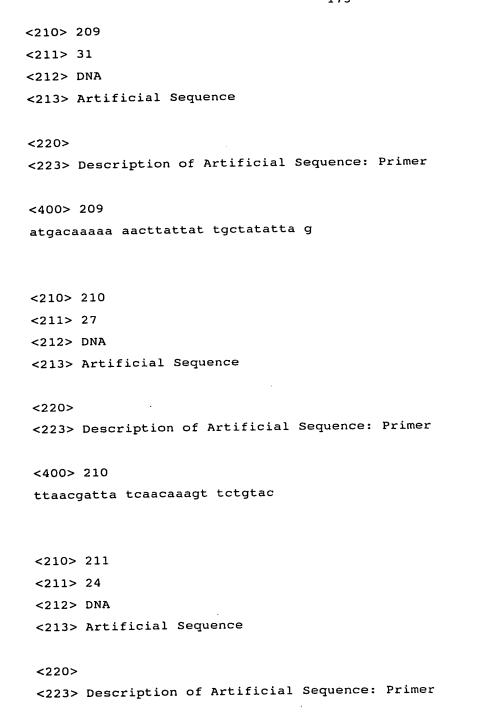


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<210> 212

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 212

ttatttatgt atagaaacag cagtccc